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FEE TRANSMITTAL for FY 2001

Patent fees are subject to annual revision.

TOTAL AMOUNT OF PAYMENT (\$ 3,302.00

Complete if Known

Application Number
Filing Date November 17, 2000
First Named Inventor Webb
Examiner Name Unknown
Group Art Unit Unknown
Attorney Docket No. TSRI 536.1Div2

METHOD OF PAYMENT

1. ☒ The Commissioner is hereby authorized to charge indicated fees and credit any overpayments to.

Deposit Account Number 19-0962
Deposit Account Name

☐ Charge Any Additional Fee Required Under 37 CFR 1.16 and 1.17

☐ Applicant claims small entity status. See 37 CFR 1.27

2. ☒ Payment Enclosed:

☒ Check ☐ Credit card ☐ Money Order ☐ Other

FEE CALCULATION

1. BASIC FILING FEE

Large Entity Fee Code (\$)	Small Entity Fee Code (\$)	Fee Description	Fee Paid
101 710	201 355	Utility filing fee	710
106 320	206 160	Design filing fee	
107 490	207 245	Plant filing fee	
108 710	208 355	Reissue filing fee	
114 150	214 75	Provisional filing fee	

SUBTOTAL (1) (\$ 710.00

2. EXTRA CLAIM FEES

Total Claims 124 -20** = 104 X Fee from below 18 = 1872
Independent Claims 12 -3** = 9 X Fee from below 80 = 720
Multiple Dependent

Large Entity Fee Code (\$)	Small Entity Fee Code (\$)	Fee Description
103 18	203 9	Claims in excess of 20
102 80	202 40	Independent claims in excess of 3
104 270	204 135	Multiple dependent claim, if not paid
109 80	209 40	** Reissue independent claims over original patent
110 18	210 9	** Reissue claims in excess of 20 and over original patent

SUBTOTAL (2) (\$ 2,592.00

**or number previously paid, if greater; For Reissues, see above

FEE CALCULATION (continued)

3. ADDITIONAL FEES

Large Entity Fee Code (\$)	Small Entity Fee Code (\$)	Fee Description	Fee Paid
105 130	205 65	Surcharge - late filing fee or oath	
127 50	227 25	Surcharge - late provisional filing fee or cover sheet	
139 130	139 130	Non-English specification	
147 2,520	147 2,520	For filing a request for <i>ex parte</i> reexamination	
112 920*	112 920*	Requesting publication of SIR prior to Examiner action	
113 1,840*	113 1,840*	Requesting publication of SIR after Examiner action	
115 110	215 55	Extension for reply within first month	
116 390	216 195	Extension for reply within second month	
117 890	217 445	Extension for reply within third month	
118 1,390	218 695	Extension for reply within fourth month	
128 1,890	228 945	Extension for reply within fifth month	
119 310	219 155	Notice of Appeal	
120 310	220 155	Filing a brief in support of an appeal	
121 270	221 135	Request for oral hearing	
138 1,510	138 1,510	Petition to institute a public use proceeding	
140 110	240 55	Petition to revive - unavoidable	
141 1,240	241 620	Petition to revive - unintentional	
142 1,240	242 620	Utility issue fee (or reissue)	
143 440	243 220	Design issue fee	
144 600	244 300	Plant issue fee	
122 130	122 130	Petitions to the Commissioner	
123 50	123 50	Processing fee under 37 CFR 1.17(q)	
126 180	126 180	Submission of Information Disclosure Stmt	
581 40	581 40	Recording each patent assignment per property (times number of properties)	
146 710	246 355	Filing a submission after final rejection (37 CFR § 1.129(a))	
149 710	249 355	For each additional invention to be examined (37 CFR § 1.129(b))	
179 710	279 355	Request for Continued Examination (RCE)	
169 900	169 900	Request for expedited examination of a design application	

Other fee (specify)

*Reduced by Basic Filing Fee Paid

SUBTOTAL (3) (\$

SUBMITTED BY

Name (Print/Type)	Emily Holmes	Registration No. (Attorney/Agent)	40,652	Complete (if applicable)	Telephone (858) 784-2937
Signature				Date	11/17/00

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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

PATENT

CERTIFICATION UNDER 37 CFR 1.10



I hereby certify that this UTILITY PATENT APPLICATION TRANSMITTAL and the documents referred to as enclosed therein are being deposited with the United States Postal Service "Express Mail Post Office to Addressee" service Mail Label No. EL587945892US under 37 CFR 1.10 on the date indicated below and is addressed to: Box Patent Application, Assistant Commissioner for Patents, Washington, D.C. 20231.

Lori A. Linderman
Lori A. Linderman

11/17/00
Date of Deposit


Applicant: Webb, et al.)	
)	
Serial No.: Unassigned)	Group Art Unit: Unassigned
)	
Filed: November 17, 2000)	Examiner: Unassigned
)	
Title: MHC CLASS II ANTIGEN- PRESENTING SYSTEMS AND METHODS FOR ACTIVATING CD4 ⁺ T CELLS)	
)	Our Ref.: TSRI 536.1Div2

PATENT

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

CERTIFICATION UNDER 37 CFR 1.10

I hereby certify that this PRELIMINARY AMENDMENT and the documents referred to as enclosed therein are being deposited with the United States Postal Service "Express Mail Post Office to Addressee" service Mail Label No. EL587945892US under 37 CFR 1.10 on the date indicated below and is addressed to: Box Patent Application, Assistant Commissioner for Patents, Washington, D.C. 20231.



Lori A. Linderman

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Applicant: Webb, et al.)	
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Serial No.: Unassigned)	Group Art Unit: Unassigned
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Filed: November 17, 2000)	Examiner: Unassigned
)	
Title: MHC CLASS II ANTIGEN- PRESENTING SYSTEMS AND METHODS FOR ACTIVATING CD4 ⁺ T CELLS)	
)	Our Ref.: TSRI 536.1Div2

PRELIMINARY AMENDMENT

Assistant Commissioner for Patents
Washington, D.C. 20231
ATTN: BOX PATENT APPLICATION

Dear Sir:

Prior to examination of the above-identified patent application on the merits, which is a Divisional application of pending prior application U.S. Serial No. 09/194,285 filed under 37 CFR 1.53(b) concurrently herewith, please enter the following amendments.

IN THE SPECIFICATION

At page 1, lines 5-6, please delete [continuation-in-part application of copending] and replace with divisional application of copending United States application Serial No. 09/194,285, filed April 12, 1999, which is based on.

IN THE CLAIMS

Please cancel claims 1-60 and 85-113.

Please add new claims 114-148.

--114. A method for activating CD4⁺ T cells in vitro, the method comprising:

a) contacting a synthetic antigen presenting cell (APC) with a peptide library in vitro for a sufficient time to generate a peptide-loaded MHC class II heterodimer for activating CD4⁺ T cells, wherein the APC comprises:

i) a MHC class II α -chain gene operably linked to a first promoter in a vector capable of expressing a MHC class II α -chain;

ii) a MHC class II β -chain gene operably linked to a second promoter in a vector capable of expressing a MHC class II β -chain, wherein upon expression of the α -chain and β -chain genes, the α -chain and β -chain form a MHC class II heterodimer capable of loading a peptide; and

iii) at least one accessory molecule gene operably linked to a third promoter in a vector capable of

expressing an accessory molecule, wherein at least one of the Class II genes and accessory molecule gene is lacking from the APC, wherein the expressed MHC class II heterodimer and accessory molecule are present on the surface of the APC in an amount sufficient for activating CD4⁺ T cells when a peptide is loaded on the MHC Class II heterodimer; and

b) contacting the peptide-loaded MHC class II heterodimer of step a) with CD4⁺ T cells, thereby inducing the contacted CD4⁺ T cells to proliferate and differentiate into activated CD4⁺ T cells.

115. The method of claim 114 further comprising:

c) separating the activated CD4⁺ T cells from the APC.

116. The method of claim 115 further comprising the step of adding the activated CD4⁺ T cells to an acceptable carrier or excipient to form a suspension.

117. The method of claim 116 further comprising the step of administering the suspension to a patient.

118. The method of claim 114 wherein the synthetic APC is produced by the method according to claim 61.

119. The method of claim 114 wherein the synthetic APC lacks a gene coding for at least one of the α -chain, the β -chain and the accessory molecule genes.

120. The method of claim 114 wherein the α - and β - chain genes are of human origin.

121. The method of claim 114 wherein the synthetic APC is an insect cell.

122. The method of claim 121 wherein the insect cell is selected from the group consisting of Spodoptera and Drosophila.

123. The method of claim 114 wherein the accessory molecule gene encodes a costimulatory molecule.

124. The method of claim 123 wherein the costimulatory molecule is B7.1 or B7.2.

125. The method of claim 114 wherein the accessory molecule gene encodes an adhesion molecule.

126. The method of claim 125 wherein the adhesion molecule is ICAM-1, ICAM-2, ICAM-3 or LFA-3.

127. The method of claim 114 wherein the accessory molecule gene encodes a survival molecule.

128. The method of claim 127 wherein the survival molecule is Fas ligand or CD70.

129. The method of claim 128 further comprising a second accessory molecule gene for a second accessory molecule.

130. The method of claim 129 wherein the first accessory molecule is a costimulatory molecule and the second accessory molecule is an adhesion molecule.

131. The method of claim 129 wherein the first accessory molecule is a costimulatory molecule and the second accessory molecule is a survival molecule.

132. The method of claim 129 wherein the first accessory molecule is a survival molecule and the second accessory molecule is an adhesion molecule.

133. The method of claim 129 further comprising a third accessory molecule gene for a third accessory molecule.

134. The method of claim 133 wherein the first accessory molecule is a costimulatory molecule, the second accessory molecule is an adhesion molecule, and the third accessory molecule is a survival molecule.

135. A method for activating CD4⁺ T cells in vitro, the method comprising:

a) contacting a cell fragment derived from the APC of a synthetic antigen presenting cell (APC) with a peptide library in vitro for a sufficient time to generate a peptide-loaded MHC class II heterodimer for activating CD4⁺ T cells, wherein the APC comprises:

i) a MHC class II α -chain gene operably linked to a first promoter in a vector capable of expressing a MHC class II α -chain;

ii) a MHC class II β -chain gene operably linked to a second promoter in a vector capable of expressing a MHC class II β -chain, wherein upon expression of the α -chain and β -chain genes, the α -chain and β -chain form a MHC class II heterodimer capable of loading a peptide; and

iii) at least one accessory molecule gene operably linked to a third promoter in a vector capable of expressing an accessory molecule, wherein at least one of the Class II genes and accessory molecule gene is lacking from the APC, wherein the expressed MHC class II heterodimer and at least one accessory molecule are operably associated on the fragment for activating CD4⁺ T cells in an amount sufficient for activating CD4⁺ T cells when a peptide is loaded on the MHC Class II heterodimer; and

b) contacting the peptide-loaded MHC class II heterodimer of step a) with CD4⁺ T cells, thereby inducing the contacted CD4⁺ T cells to proliferate and differentiate into activated CD4⁺ T cells.

136. The method of claim 135 further comprising:

c) separating the activated CD4⁺ T cells from the APC.

137. The method of claim 136 further comprising the step of adding the activated CD4⁺ T cells to an acceptable carrier or excipient to form a suspension.

138. The method of claim 137 further comprising the step of administering the suspension to a patient.

139. The method of claim 135 wherein the cell fragment is derived from a synthetic APC produced by the method according to claim 61.

140. A method of altering a CD4⁺ T cell-mediated immune response to treat a condition in a patient comprising:

- a) analyzing the patient for patient-specific cytokine profile;
- b) collecting CD4⁺ T cells from the patient;
- c) contacting the CD4⁺ T cells with a synthetic antigen presenting cell (APC) for a sufficient time to activate CD4⁺ T cells, wherein the APC comprises:
 - i) a MHC class II α -chain gene operably linked to a first promoter in a vector capable of expressing a MHC class II α -chain;
 - ii) a MHC class II β -chain gene operably linked to a second promoter in a vector capable of expressing a MHC class II β -chain, wherein upon expression of the α -chain and β -chain genes, the α -chain and β -chain form a MHC class II heterodimer capable of loading a peptide; and
 - iii) at least one accessory molecule gene operably linked to a third promoter in a vector capable of expressing an accessory molecule, wherein at least one of the Class II genes and accessory molecule gene is lacking from the APC, wherein the expressed MHC class II heterodimer and accessory molecule are present on

the surface of the APC in a sufficient amount to activate CD4⁺ T cells when a peptide is loaded onto the heterodimer in vitro thereby inducing the contacted CD4⁺ T cells to proliferate and differentiate into activated CD4⁺ T cells that produce a functionally opposing cytokine profile to the profile obtained in step a); and

d) returning the activated CD4⁺ T-cells to the patient.

141. The method of claim 140 wherein the condition is an autoimmune disease.

142. The method of claim 141 wherein the autoimmune disease is selected from the group consisting of diabetes, multiple sclerosis, autoimmune thyroiditis, systemic lupus erythematosus, myasthenia gravis, Crohn's disease and inflammatory bowel disease.

143. The method of claim 141 wherein the patient-specific cytokine profile is produced by a CD4⁺ Th1 type response.

144. The method of claim 143 wherein the patient-specific cytokine profile comprises the cytokine selected from the group consisting of interleukin-2, interferon- γ and tumor necrosis factor.

145. The method of claim 140 wherein the condition is an allergy.

146. The method of claim 145 wherein the allergy is selected from the group consisting of asthma and contact sensitivity.

147. The method of claim 145 wherein the patient-specific cytokine profile is produced by a CD4⁺ Th2 type response.

148. The method of claim 147 wherein the patient-specific cytokine profile comprises the cytokine selected from the group consisting of interleukin-4 and interleukin-10. --

REMARKS

Claims 1-60 and 85-113 have been cancelled and new claims 114-148 have been added. Claims 61-84 and 114-148 are now pending. No new matter has been added.

Respectfully submitted,

11/17/00
Date

Emily Holmes
Emily Holmes, Reg. No. 40,652

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[] Filed Under §1.34(a)

MHC CLASS II ANTIGEN PRESENTING SYSTEMS AND
METHODS FOR ACTIVATING CD4⁺ T CELLS

Cross-Reference to Related Applications

5 This application is a continuation-in-part application of
copingding United States provisional application Serial No.
60/018,175, filed May 23, 1996, having the same title as above,
the disclosure of which are hereby incorporated by reference.

10 Technical Field

The present invention relates to materials and methods of
activating CD4⁺ T cells with specificity for particular antigenic
peptides, the use of activated T cells in vivo for the treatment
of a variety of disease conditions, and compositions appropriate
15 for these uses.

Background

Though the T cell repertoire is largely shaped during T
cell development in the thymus, mature CD4⁺ T cells are also
20 regulated extrathymically. Whereas, some conditions of
activation lead to tolerance reflecting either anergy or clonal
elimination, other conditions lead to a change in the type of
response observed. For CD4⁺ T cells, this change in functional
phenotype is largely a change in the pattern of cytokines
25 produced. Although CD4⁺ T cells that are subject to acute
activation maintain the ability to produce multiple cytokines, T
cells obtained under conditions of chronic stimulation
frequently demonstrate a more restricted pattern of cytokine
production. For example, T cell clones maintained by repeated
30 stimulation in vitro have defined two major functional
categories of CD4⁺ T cells referred to as Th1 and Th2 type cells.
Th1 type cells produce primarily interleukin-2 (IL-2),
interferon- γ (IFN- γ) and tumor necrosis factor (TNF), all of

which are referred to as inflammatory cytokines. In contrast, Th2 type cells typically produce IL-4, IL-5, and IL-10 and are important for antibody production and for regulating the responses of Th1 type cells.

5 Although such extreme segregation in cytokine production is often not seen during in vivo T cell responses, recovery from certain types of infections, such as Leishmania, is associated with preferential production of IL-2/IFN- γ . Mice that mount a Th2 response to Leishmania fail to contain the infection and
10 ultimately die. Inappropriate production of cytokines of the Th2 type response has been frequently linked to allergic type diseases such as asthma and contact sensitivity. For review on activation of CD4⁺ T cells and role in allergic disease, see Hetzel and Lamb, Clinical Immunol. Immunopath., 73:1-10 (1994).

15 Perhaps the strongest association of human disease with skewed patterns of cytokine production is the association of Th1 responses and Th1 type cytokines with autoimmune disease. Strong evidence in experimental models indicates that many types
20 of autoimmunity including diabetes, experimental models for multiple sclerosis, autoimmune thyroiditis, and the like are mediated by Th1 type CD4⁺ T cells. The expression of Th2-associated cytokines, such as IL-4, in these models interfere with the development of autoimmune disease. Th2 type cytokines
25 dampen the response of Th1 type cells while the Th1 type cytokines antagonize the development of Th2 type responses.

 In view of the association of particular activated T cell subsets with particular disease conditions, a need therefore exists to be able to direct the proliferation and activation of
30 CD4⁺ T cells to a desired T cells subset, a process that is extremely beneficial in altering the course of disease. One potential solution is to activate in vitro CD4⁺ T cells that are first isolated from a subject who may optionally be having

either allergy or autoimmune conditions to produce cells secreting a preferred cytokine profile. The resultant activated T cells are then reintroduced to the subject to alter the course of disease and perhaps even provide a long term cure.

5 The challenge in this approach, now solved by the present invention, is the difficulty in defining activation conditions that reproducibly generate CD4⁺ T cell subsets that produce the desired therapeutic cytokine profile. Expression of particular cytokines is linked to a particular antigen presenting cell
10 (APC) and their associated accessory (assisting) molecules. For a review of the surface proteins serving as accessory molecules that are involved in T cell costimulation, see Mondino and Jenkins, J. Leukocyte Biol., 55:805-815 (1994). Since both the cytokines produced by the APC and the coordinately expressed
15 accessory molecules are themselves regulated by multiple factors, including the type of antigen, the affinity of the T cell receptor (TCR)-antigen interaction, antigen concentration and the like, predicting the outcome of T cell activation upon antigen presentation is historically very difficult. Indeed, as
20 additional accessory molecules have been proposed for the activation process in vivo, it has become increasingly clear that many diverse molecules are involved in the regulation of T cell responses and act in combinatorial fashion to effect the outcome of T cell activation.

25 Prior to the present invention, the co-expression of selected MHC class II molecules in conjunction with one or more selected accessory molecules has not been possible. The present invention now presents a solution to predictably generate a preferred T cell phenotype through the reproducible activation
30 of T cells to generate either Th1 or Th2 type T cells. The invention describes the generation of synthetic APC that present, in a neutral background, MHC class II molecules in combination with defined accessory molecules. The MHC class II

molecules and defined accessory molecules are expressed in a nonmammalian insect cell and can be presented in a variety of forms of synthetic APC including insect cells displaying the molecules.

5 The advantage of using the insect cells as the expression and presentation vehicles for the MHC class II/accessory molecule compositions of this invention is that the cells do not endogenously produce regulatory cytokines and do not express mammalian accessory molecules. This overcomes the inherent
10 unpredictability of using mammalian APC that express many molecules that are capable of altering the T cell response. In addition, the insect cell expression system described in the present invention provides for the expression of MHC class II molecules without bound peptide (i.e., "empty" molecules) that
15 can be produced under certain restrictive circumstances, such as temperature requirements. At physiological temperatures, these "empty" molecules are normally unable to reach the cell surface as class II without bound peptide are very thermolabile. The invention utilizes the capacity of "empty" MHC class II
20 compositions to allow for the exogenous loading of selected peptides along with the ability to provided endogenously loaded counterparts.

 A recombinant glycosyl-phosphatidylinositol (GPI)-modified MHC class I molecule (HLA-A2.1:GPI/ β_2m) was generated in the
25 above-described insect cell system to produce antigen presenting cells as described in International Publication Number WO 96/12009 by Tykocinski. In that publication, the recombinant GPI-modified MHC class I molecules are isolated from the insect cell by affinity purification for subsequent reincorporation
30 into cell membranes. In other aspects, the publication describes the preparation of a GPI-modified MHC class I molecule co-anchored on a cell membrane with a GPI-modified B7.1 costimulatory molecule. Although the publication states that

GPI-modified MHC class II molecules can be prepared as described for those of MHC class I, the publication does not present any details for such preparation.

In contrast, the present invention provides and describes a unique means based on the co-expression of a selected MHC class II haplotype in conjunction with one or more accessory molecules, such as B7.1, to activate CD4⁺ T cells resulting in the differentiation to a particular T cell subset, Th1 or Th2 cells, that effect a preferred cytokine profile influence. The invention provides the advantage of selectively activating CD4⁺ T cells in vitro to a preferred T cell subset thereafter allowing for the reintroduction of the activated T cells into the patient. The present invention thus provides the ability to combine individual presenting molecules with particular accessory molecules for expression in selected combinations that permits reproducibility and predictability for selectively activating CD4⁺ T cells to a desired T cell subset not available in other approaches.

Brief Summary of the Invention

It has now been discovered that recombinant MHC class II molecules expressed in combination with selected accessory molecules, including costimulatory molecules and adhesion molecules, are effective in activating CD4⁺ T cells to become armed effector T cells that recognize target cells on which MHC class II heterodimer is expressed for complexation with peptide. Activation is characterized by proliferation and differentiation into effector T cell subsets, Th1 and Th2, that secrete particular cytokines. Th1 and Th2 type T cells are respectively referred to as inflammatory cells and T-helper cells.

Thus, the present invention relates to a synthetic antigen presenting system, also referred to as APC, for producing and presenting a mammalian, preferably human, MHC class II molecule

in combination with one or more accessory molecules to activate CD4⁺ T cells.

In one embodiment, the system relates to a synthetic antigen presenting matrix having a support and at least the
5 extracellular portion of a MHC class II heterodimeric molecule operably linked to the support and capable of binding to a selected peptide. The matrix also includes an accessory molecule operably linked to the support. The accessory molecule interacts with a specific receptor present on the CD4⁺ T cell.
10 The MHC class II and accessory molecules are present in sufficient numbers to activate a population of CD4⁺ T cells specific for the MHC class II/peptide combination when the peptide is bound to the extracellular portion of the MHC molecule.

15 It has been found that an antigen presenting matrix having both a MHC class II heterodimer or at least the extracellular portion thereof loaded with a peptide specific for that MHC, together with an accessory molecule, provides a synergistic reaction in activating CD4⁺ T cells. Examples of accessory
20 molecules are costimulatory molecules, including B7.1 and B7.2, adhesion molecules such as intercellular cell adhesion molecule-1 (ICAM-1) and lymphocyte function-associated antigen-3 (LFA-3), and survival molecules such as Fas ligand (FasL) and CD70. In some embodiments, the extracellular portion of such accessory
25 molecules can also be used in the present invention.

The support used for the matrix can take several different forms. Examples for the support include solid support such as metals or plastics, porous materials such as resin or modified cellulose columns, microbeads, microtiter plates, red blood
30 cells and liposomes.

Another type of support is a cell fragment, such as a cell membrane fragment. An entire cell is also contemplated as a support. In this embodiment, the matrix is actually cells which

have been transformed with one or more expression vectors containing genes for the expression of MHC class II α - and β -chains along with at least one accessory molecule. The expressed proteins are then transported to the cell membrane where the transmembrane domain of the class II chains provide anchors allowing the extracellular domain to be displayed on the outer cell surface, thereby creating a synthetic antigen presenting cell (APC). The expression vectors contain the selected genes, preferably in the form of a cDNA sequence, operably linked to a promoter that is either constitutive or inducible.

The MHC α - and β -chains associate together forming a MHC class II heterodimer which binds to a peptide specific for that heterodimer. With the present invention, two methods of loading peptides onto MHC class II heterodimers are contemplated. In one embodiment, the peptide is loaded intracellularly following proteolytic processing of internalized intact protein into peptide fragments. The peptides are then loaded onto newly generated MHC class II molecules while they are still within the cell. Alternatively, the MHC class II molecules are expressed as empty molecules on the cell surface and synthetic or processed peptide reagents are then loaded extracellularly onto the MHC class II heterodimer.

Nucleotide sequences for encoding at least one accessory molecule gene operably linked to a promoter in a vector are also introduced into the cell. Following expression, the accessory molecule is coordinately anchored on the surface of the cell along with the MHC class II heterodimer in sufficient numbers to activate a population of CD4⁺ T cells lymphocytes specific for the MHC class II/peptide complex. Other molecules referred to as antigen processing assisting molecules are also contemplated for use in generating recombinant APC. These molecules are either provided by the cell used as APC or exogenously through

an expression vector system as described above. Examples of such antigen processing assisting molecules include invariant chain, lysosomal enzymes and H2-M and H2-O molecules.

The cell line is synthetic in that at least one of the
5 genes described above is not naturally present in the cells from which the cell line is derived. It is preferable to use a poikilotherm cell line because MHC molecules are thermolabile. A range of species are useful for this purpose. See, for example, U.S. Patent No. 5,314,813 to Petersen et al. which
10 discusses numerous species for this use, the disclosure of which is hereby incorporated by reference. Eukaryotic cells and preferably insect cells are used as APC. Preferred insect cells include Drosophila (fruit fly) and Spodoptera (butterfly).

MHC class II molecules have been expressed in insect cells
15 such as Drosophila and Spodoptera cells. Since these cells do not have all the components of a mammalian immune system, the various proteins involved in the peptide loading machinery are absent from such cells. The lack of mammalian peptide-loading machinery allows the introduced mammalian MHC class II molecules
20 to be expressed as empty molecules at the cell surface when the cells are cultured at thermostabile temperature restrictive conditions, such as at 28°C. In contrast, at 37°C, empty Class I molecules are thermolabile and tend to disintegrate. Thus, by incubating MHC class II-expressing Drosophila cells with
25 peptides that specifically bind to anchored MHC class II molecule, virtually every class II molecule is loaded with one and the same peptide. Moreover, the invention provides for the means to introduce any known MHC class II α - and β -chain genes into an expression vector thereby overcoming the inherent limit
30 to the number of MHC class II molecules expressed in any one mammal.

In the present invention, a specifically effective synergistic reaction in driving CD4⁺ T cells to a Th1-type

response characterized by an increase in the cytokine interleukin-2 (IL-2) results from a Drosophila antigen presenting cell having MHC class II molecules bound with a peptide, a costimulatory molecule, and an adhesion molecule. In particular, a highly effective synergistic generation of IL-2 production coupled with CD4⁺ proliferation results from the combination of B7.2 and ICAM-1. In contrast, without ICAM-1 but with either B7.1 or B7.2, the Drosophila APC system loaded with peptide induced a Th2-type response characterized by an increase in IL-4. Thus, ICAM-1 antagonized the Th2-type response resulting in a Th1-type phenotype.

A Th1 phenotype characterized by IL-2 production coupled with proliferative responses also resulted from a synthetic antigen presenting cell having CD70 expressed simultaneously with ICAM-1 with or without B7.2.

Therefore, the selection of MHC class II genes in combination with at least one accessory molecule genes for expression thereof in an APC of this invention can be tailored depending upon the desired outcome for effecting proliferation and phenotypic activation of CD4⁺ T cells.

The present invention also relates to methods for making the synthetic APC systems as described above in which at least one expression vector containing genes for a MHC class II heterodimer and an accessory molecule is introduced.

Methods of producing activated CD4⁺ T cells in vitro are also contemplated. One preferred method comprises contacting, in vitro, CD4⁺ cells with a synthetic MHC class II/accessory molecule-bearing APC described above for a time period sufficient to activate, in an antigen-specific manner, a population of CD4⁺ T cells. The method may further comprise (1) separating the activated CD4⁺ cells from the antigen-presenting matrix; (2) suspending the activated CD4⁺ cells in an acceptable carrier or excipient; and (3) administering the suspension to an

individual in need of treatment. As previously discussed, the antigens may comprise native or undegraded proteins or polypeptides, or they may comprise antigenic polypeptides which have been cleaved or synthesized into peptide fragments comprising at least 8 amino acid residues prior to incubation with the mammalian MHC class II heterodimeric molecules.

In addition to the utility of being able to direct the activation of CD4⁺ T cells to a desired T cell subset as described above, the ability to express any MHC class II molecule provides the means to identify unknown CD4⁺-activating peptide specific for that particular MHC class II molecule. As such, the present invention contemplates the activation of CD4⁺ T cells through the screening of a peptide library with synthetic APC expressing a particular MHC class II heterodimer.

In a further embodiment, the synthetic APC system described herein is useful for isolation of reactive CD4⁺ T cells from a heterologous population of cells. Such isolation provides the ability to monitor ongoing CD4⁺ T cell-mediated responses in disease conditions in a patient.

In another variation of the above, in view of the ability to selectively activate CD4⁺ T cells into a particular T cell subset for producing a preferred cytokine profile, the invention relates to methods of treating conditions in patients mediated by a undesirable CD4⁺ response. Such disease conditions characterized by either a Th1- or Th2-type response include autoimmune diseases, allergy and cancer. The therapeutic goal is to introduce CD4⁺ T cells activated to a preferred T cell subset to antagonize an ongoing CD4⁺ T cell response. Thus, the method comprises (1) obtaining a fluid sample containing resting or naive CD4⁺ cells from the patient; (2) contacting, in vitro, the CD4⁺ cells with a selected synthetic peptide-loaded APC of this invention for a time period sufficient to activate, in an

antigen-specific manner, the CD4⁺ cells; and (3) administering the activated CD4⁺ cells to the patient.

Other embodiments are apparent to one skilled in the art.

5 Brief Description of the Drawings

Figures 1A-1C diagram the construction of expression plasmids pRmHa-2 and pRmHa-3. In Figure 1A, pRmHa-2 construction is shown; in Figure 1B, pRmHa-3 construction is shown; and in Figure 1C, the pRmHa-3 vector is illustrated, showing the restriction, polylinker, promoter, and polyadenylation sites, as well as a site at which a nucleotide sequence may be inserted for expression.

Figure 2 shows the proliferative response of DO10 T cell receptor (TCR) transgenic mouse cell line (Tg cells) when cultured in the presence of Drosophila cell lines with or without ovalbumin (OVA) peptide loaded onto the surface-expressed MHC class II heterodimer. Proliferation is assayed as described in Example 5. The proliferation is measured in counts per minute (cpm) X 1000 as plotted on the Y-axis against increasing concentrations of OVA peptide in μ M on the X-axis. T-S (diamond marked line) shows responses with control splenic APC. Proliferative responses with recombinant MHC class II alone are shown in the line having closed circles. Those with MHC class II combined with either costimulatory molecules B7.1 or B7.2 are shown respectively with lines having open and closed squares.

Figure 3 shows proliferative responses to recombinant MHC class II alone (closed circle line), with MHC class II plus B7.2 (closed square line), with MHC class II plus ICAM-1 (open triangle line) and with MHC class II plus B7.2 and ICAM-1 (open diamond line). Refer to Figure 2 legend for other details.

Figures 4A-4D show the cytokine profile produced in response to activation of CD4⁺ T cells when cultured in the

presence of Drosophila APC having recombinant MHC class II alone or in combination with B7.1 or B7.2 costimulatory molecules. Splenic APC (labeled T-S) are control assays. The assays are performed as described in Example 5. Figures 4A-4D respectively show the cytokines Il-2, Il-4, IFN- γ and Il-10 in ng/ml as plotted on the Y-axis. Cytokine profiles were assessed over three days between day 3 and 5 of culture.

Figures 5A-5D show the cytokine profile produced in response to activation of CD4⁺ T cells when cultured in the presence of Drosophila APC having recombinant MHC class II alone or in combination with B7.2 costimulatory molecule, ICAM-1 and with B7.2 and ICAM-1. Figures 5A-5D respectively show the cytokines Il-2, Il-4, IFN- γ and Il-10 in ng/ml as plotted on the Y-axis. Refer to Figure 4 legend for other details.

Detailed Description of the Invention

A. Definitions

Amino Acid Residue: An amino acid formed upon chemical digestion (hydrolysis) of a polypeptide at its peptide linkages. The amino acid residues described herein are preferably in the "L" isomeric form. However, residues in the "D" isomeric form can be substituted for any L-amino acid residue, as long as the desired functional property is retained by the polypeptide. NH₂ refers to the free amino group present at the amino terminus of a polypeptide. COOH refers to the free carboxy group present at the carboxy terminus of a polypeptide. In keeping with standard polypeptide nomenclature (described in J. Biol. Chem., 243:3552-3559 (1969) and adopted at 37 CFR §1.822(b)(2)), abbreviations for amino acid residues are shown in the following Table of Correspondence:

TABLE OF CORRESPONDENCE

<u>SYMBOL</u>		<u>AMINO ACID</u>
<u>1-Letter</u>	<u>3-Letter</u>	
Y	Tyr	tyrosine
5 G	Gly	glycine
F	Phe	phenylalanine
M	Met	methionine
A	Ala	alanine
S	Ser	serine
10 I	Ile	isoleucine
L	Leu	leucine
T	Thr	threonine
V	Val	valine
P	Pro	proline
15 K	Lys	lysine
H	His	histidine
Q	Gln	glutamine
E	Glu	glutamic acid
Z	Glx	Glu and/or Gln
20 W	Trp	tryptophan
R	Arg	arginine
D	Asp	aspartic acid
N	Asn	asparagine
B	Asx	Asn and/or Asp
25 C	Cys	cysteine
X	Xaa	Unknown or other

It should be noted that all amino acid residue sequences represented herein by formulae have a left- to-right orientation in the conventional direction of amino terminus to carboxy terminus. In addition, the phrase "amino acid residue" is broadly defined to include the amino acids listed in the Table of Correspondence and modified and unusual amino acids, such as

those listed in 37 CFR 1.822(b)(4), and incorporated herein by reference. Furthermore, it should be noted that a dash at the beginning or end of an amino acid residue sequence indicates a peptide bond to a further sequence of one or more amino acid
5 residues or a covalent bond to an amino-terminal group such as NH_2 or acetyl or to a carboxy-terminal group such as COOH .

Recombinant DNA (rDNA) molecule: A DNA molecule produced by operatively linking two DNA segments. Thus, a recombinant DNA molecule is a hybrid DNA molecule comprising at least two
10 nucleotide sequences not normally found together in nature. rDNA's not having a common biological origin, i.e., evolutionarily different, are said to be "heterologous".

Vector: A rDNA molecule capable of autonomous replication in a cell and to which a DNA segment, e.g., gene or
15 polynucleotide, can be operatively linked so as to bring about replication of the attached segment. Vectors capable of directing the expression of genes encoding for one or more polypeptides are referred to herein as "expression vectors".

Upstream: In the direction opposite to the direction of
20 DNA transcription, and therefore going from 5' to 3' on the non-coding strand, or 3' to 5' on the mRNA.

Downstream: Further along a DNA sequence in the direction of sequence transcription or read out, that is traveling in a 3'- to 5'-direction along the non-coding strand of the DNA or
25 5'- to 3'-direction along the RNA transcript.

Reading Frame: A particular sequence of contiguous nucleotide triplets (codons) employed in translation that define the structural protein encoding-portion of a gene, or structural gene. The reading frame depends on the location of the
30 translation initiation codon.

Polypeptide: A linear series of amino acid residues connected to one another by peptide bonds between the alpha-amino group and carboxy group of contiguous amino acid residues.

Protein: A linear series of greater than 50 amino acid residues connected one to the other as in a polypeptide.

Receptor: A receptor is a molecule, such as a protein, glycoprotein and the like, that can specifically (non-randomly) bind to another molecule.

Substantially Purified or Isolated: When used in the context of polypeptides or proteins, the terms describe those molecules that have been separated from components that naturally accompany them. Typically, a monomeric protein is substantially pure when at least about 60% to 75% of a sample exhibits a single polypeptide backbone. Minor variants or chemical modifications typically share the same polypeptide sequence. A substantially purified protein will typically comprise over about 85% to 90% of a protein sample, more usually about 95%, and preferably will be over about 99% pure. Protein or polypeptide purity or homogeneity may be indicated by a number of means well known in the art, such as polyacrylamide gel electrophoresis of a sample, followed by visualization thereof by staining. For certain purposes, high resolution is needed and high performance liquid chromatography (HPLC) or a similar means for purification utilized.

Synthetic Peptide: A chemically produced chain of amino acid residues linked together by peptide bonds that is free of naturally occurring proteins and fragments thereof.

25

B. MHC Class II Heterodimers, Accessory Molecules and Antigen Processing Assisting Molecules

The present invention relates to a synthetic antigen-presenting system for use in activating CD4⁺ T cells. Though the T cell repertoire is largely shaped during T cell development in the thymus, mature CD4⁺ T cells are also regulated extrathymically. Although some conditions of activation lead to tolerance reflecting either anergy or clonal elimination, other

conditions lead to a change in the type of response observed. For CD4⁺ T cells, this change in functional phenotype is largely a change in the pattern of cytokines produced. Although CD4⁺ T cells that are subject to acute activation maintain the ability to produce multiple cytokines, T cells obtained under conditions of chronic stimulation frequently demonstrate a more restricted pattern of cytokine production. For example, T cell clones maintained by repeated stimulation in vitro have defined two major functional categories of CD4⁺ T cells referred to as Th1 and Th2 type cells. Th1 type cells produce primarily interleukin-2 (IL-2), interferon- γ (IFN- γ) and tumor necrosis factor (TNF), all of which are referred to as inflammatory cytokines. As such, the TH1 cells are sometimes referred to as inflammatory T cells which then activate macrophages to kill intravesicular pathogens they harbor as mediated by peptides generated from processing of pathogenic proteins presented to CD4⁺ T cells. In contrast, Th2 type cells typically produce IL-4, IL-5, and IL-10 and are important for antibody production and for regulating the responses of Th1 type cells. As such, Th2 cells are sometimes referred to as helper T cells that activate B cells to make antibody in response to peptides derived from extracellular pathogens and toxins that are presented to CD4⁺ T cells.

A CD4⁺ T cell is defined as a T cell or a T lymphocyte that has the CD4⁺ co-receptor on its cell surface in conjunction with the presence of either $\alpha:\beta$ or $\gamma:\delta$ heterodimeric receptor associated with the proteins of the CD3 complex.

Activation of CD4⁺ T cell subsets is characterized by proliferation of the responsive T cell population coordinated with the selection production of cytokines as described above dependent upon the type of stimulation induced by an antigen presenting cell. The latter is defined as a highly specialized cell that can process antigens and display their peptide

fragments on the cell surface together with molecules required for lymphocyte activation. The specificity of CD4⁺ T cell activation is based on the T cell antigen receptor (TCR) recognition of peptide antigens bound to MHC class II

5 heterodimers on the surface of antigen presenting cells (APC). The main APC for T cells are dendritic cells, macrophages and B cells. In addition, APC-derived non-antigenic costimulatory signals play a contributory role in CD4⁺ T cell activation.

The present invention utilizes a synthetic antigen
10 presenting system, based on the natural immunological response mechanisms described above, to manipulate the activation of CD4⁺ T cells by recombinant MHC class II molecules in conjunction with one or more accessory molecules broadly referred to as costimulatory molecules. The latter include specific
15 costimulatory molecules, adhesion molecules and survival molecules, and the like. In other aspects of the invention, the synthetic antigen presenting system further contains antigen processing assisting molecules that are useful in generating peptide-loaded MHC class II molecules. In the context of this
20 invention, the synthetic antigen presenting systems are useful for activating CD4⁺ T cells in vitro and in vivo. These aspects are discussed in Section E on Methods of Altering CD4⁺ T Cell Responses.

Thus, as mentioned above, the synthetic antigen presenting
25 system of the present invention has at least two major components. The first component is at least the extracellular portions of a recombinant MHC class II heterodimer which is capable of binding to a peptide that provides the specificity of CD4⁺ T cell activation via recognition by the TCR. The second
30 major component is at least the extracellular portion of at least one accessory molecule that provides a non-antigenic specific costimulatory signal in the activation of CD4⁺ T cells. In other embodiments, the entire molecules of the antigen

presenting system and the non-antigen costimulatory signals are used.

For ease of description, MHC class II heterodimers will be discussed generally, with the understanding that an
5 extracellular portion of the MHC molecule may be used in certain aspects of the invention. The portion of the MHC molecule necessary for the present invention is the part that binds to the antigenic peptide for presentation to the CD4⁺ T cells.

The present invention allows the recombinant MHC class II
10 heterodimers to be produced by vector-transformed synthetic antigen presenting cells with the peptide already complexed with the MHC class II heterodimer. Alternatively, empty MHC class II heterodimers are produced that do not yet have a peptide complexed with them. This latter embodiment is particularly
15 useful as it allows for complexation with a particular peptide or for screening of a library of peptides after the MHC class II heterodimers are expressed.

1. MHC Class II Genes and Encoded Heterodimers

20 MHC class II molecules are cell surface glycoproteins that consist of a non-covalent complex of two chains, α and β , both of which span the membrane. A peptide-binding cleft is formed between the cooperating chains. Peptides that bind MHC class II molecules are variable in length and have anchor
25 residues that lie at various distances from the ends of the peptides thereby resulting in peptides having ends that are not tightly bound within the cleft of the binding pocket. See, Janeway and Travers, Immunobiology, Section 4-4, Current Biology LTD, 2nd ed., 1996. Further aspects of the presenting antigenic
30 peptides are discussed in Section D.

In vivo, empty MHC class II heterodimers become destabilized and are subsequently removed from the cell surface thereby preventing MHC molecules for acquiring peptides from the

surrounding extracellular fluid that would deleteriously effect T cell specificity. In present invention, the synthetic antigen presenting cell system allows for the production of empty MHC class II heterodimers on the cell surface that are not subject to destabilizing events. As a consequence, loading of an antigenic peptide on a surface-expressed recombinant MHC class II heterodimer is facilitated for subsequent use in manipulating CD4⁺ activation and cytokine production.

The present invention, particular combinations of a selected MHC class II molecule with a coordinate antigen is facilitated by the presence of consensus nucleotide sequences in MHC class II genes. These regions, described further below, allow for the retrieval and use of the multiple MHC class II genes in the MHC complex in mammals as well as the multiple alleles of each gene. In other words, MHC is both polygenic and polymorphic having respectively several genes and multiple alleles of each gene.

In humans, MHC is called HLA while in mouse it is referred to as H-2. Three pairs of MHC class II α - and β -chain genes in humans have been designated HLA-DP, HLA-DQ and HLA-DR. The HLA-DR cluster contains an extra β -chain gene. As such, the three sets of genes can give rise to four types of MHC class II molecules. MHC class II genes and encoded α - and β -chains that are obtained from human genes are said to be of human origin. In mice, MHC class genes are designated H2-M, H2-A and H2-E. As each MHC class II molecule binds a different range of peptides, the presence of multiple gene loci gives an individual the ability to present a broad range of different peptides than if only one MHC class II molecules of each class were expressed at the cell surface.

While the polymorphic MHC class II genes encode corresponding proteins that vary by only one or a few amino acids, the different allelic variants differ by up to 20 amino

acids. As a result, MHC class II diversity expands the ability of antigen recognition by T cells. Moreover, via MHC restriction, T cells have been shown to recognize peptide in the context of a particular MHC molecule but not when presented on another. Thus, T cell receptor specificity is imparted by both peptide and by the MHC molecule binding it.

The MHC class II heterodimers of this invention containing selected α - and β -chains are obtained by amplification of MHC class II-encoding genes and allelic variants thereof with specific pairs of oligonucleotide primers. The nucleotide sequences of the primers allow for amplification of the diversity of MHC class II genes and allelic variants thereof based on the 5' and 3' consensus nucleotide sequences present in the genes within a category of genes. Specific nucleotide sequences of primer pairs for amplifying the α - and β -chains of human HLA-DP, -DQ and -DR genes as well as those for amplifying murine IA^d-encoding heterodimers are presented in Example 2A.

The MHC class II-encoding genes are amplifiable from a variety of cellular sources including B cells, macrophages and dendritic cells, all of which are present in the blood. The amplification conditions for obtaining amplified MHC class II-encoding genes with the primers of this invention are described in Section C.

The α - and β -chains comprising the MHC heterodimers of this invention are useful in either anchored or soluble form. In the anchored form, the recombinant MHC heterodimer is anchored into the synthetic antigen presenting cell from which it is expressed. Alternatively, a recombinant MHC heterodimer is anchored in a matrix comprising a support as described in this invention after being secreted in soluble form. The latter is generated when a stop codon is engineered during the amplification procedure or thereafter into the nucleotide

sequence encoding the MHC class II α - and β -chains of choice preceding the transmembrane domain.

2. Accessory Genes and Encoded Molecules

5 The accessory molecules of this invention, including costimulatory molecules, adhesion molecules and survival molecules, are effective in concert with the MHC class II heterodimer complexed with peptide in activating CD4⁺ T cells to become armed effector T cells that recognize target cells.

10 Naive T cells are activated to proliferate and differentiate into armed effector T cells when they encounter their specific antigen when presented by a peptide-loaded MHC class II heterodimer on the surface of an APC. Activation not only requires the recognition of a foreign peptide fragment bound to

15 a MHC class II heterodimer but it also requires the simultaneous delivery of a costimulatory signal concurrently expressed by the APC.

Thus, the synthetic APC or matrices of this invention are characterized by the presence not only of a particular MHC class

20 II heterodimer but also by the presence of one or more costimulatory molecules that are broadly defined as accessory molecules. At least three types of accessory molecules, including specific costimulatory molecules, adhesion molecules, and survival molecules, are contemplated for use in preparing

25 synthetic APC or matrices of this invention

a. Costimulatory Molecules

A first type of an accessory molecule is composed of costimulatory molecules such as B7.1 (previously known as B7

30 and also known as CD80) and B7.2 (also known as CD86) which binds to CD28 on T cells. B7.1 and B7.2 are structurally related glycoproteins that are homodimeric members of the immunoglobulin superfamily. Other costimulatory molecules are

anti-CD28 antibodies or the functional portions of such antibodies, e.g. Fab portions that bind to CD28. Ligation of CD28 by the above molecules has been shown to costimulate the growth of naive T cells. On activated T cells, an additional
5 receptor, CTLA-4, binds B7 molecules with a higher affinity than that with CD28.

Recombinant B7 costimulatory molecules for use in the synthetic APC or matrices of this invention are obtained by PCR as described for MHC class II molecules. Preferred
10 oligonucleotide primers and cellular sources for amplification therefrom as described in Example 2C.

b. Adhesion Molecules

Another major type of accessory molecule of the
15 present invention is an adhesion molecule that also functions in T cell activation. Accessory adhesion molecules include the various ICAM molecules, which include intercellular adhesion molecule (ICAM) ICAM-1, ICAM-2, ICAM-3, lymphocyte function-associated antigen (LFA) LFA-1 and LFA-3. All of these
20 molecules are members of the immunoglobulin superfamily. The ICAM-related members all bind to the T cell integrin, LFA-1. In addition to being expressed on APC including dendritic cells, macrophages and B cells, ICAM-1 and ICAM-2 are also expressed on endothelium, thereby mediating cell adhesion and subsequent
25 extravasation between circulating leukocytes and endothelium. ICAM-3, however, is only expressed on leukocytes and is thought to play an important part in adhesion between T cells and APC.

The interaction between ICAM-1, -2 and -3 synergizes with a second adhesive interaction between LFA-3 (CD58) and LFA-2 (CD2)
30 that are respectively expressed on an APC and a T cell surface.

Recombinant adhesion molecules for use in the synthetic APC or matrices of this invention are obtained by PCR as described

for MHC class II molecules. Preferred oligonucleotide primers and cellular sources for amplification therefrom as described in Example 2C.

5 c. Survival Molecules

A survival molecule is another type of an accessory molecule that plays a role in metabolic responses ranging from stimulatory to inducing cell death. Thus, a survival molecule can also be referred to as a cell death
10 regulating molecule. A survival molecule is typically a protein but may include other types of macromolecules such as carbohydrates, lipids and the like. Survival molecules for use in the compositions and methods of this invention include Fas ligand, TNF-receptor, TNF, CD70, a Type II transmembrane protein
15 that is a member of the TNF family that binds to CD27, a member of the TNF receptor family. Fas ligand binds to the receptor called Fas and receptor occupancy resulting in the induction of apoptotic cell death of the cell expressing Fas receptor. CD27 is expressed on resting T and B cells while CD70 is expressed on
20 activated T and B cells. Binding of CD70 to its receptor, CD27, induces T-cell costimulation and the interaction may be important for the recruitment of T cells from the unprimed T cell pool. Under other certain conditions, activation of the TNF receptor by TNF results in a similar response.

25 The recombinant survival molecules described above for use in the synthetic APC or matrices of this invention are obtained by PCR as described for MHC class II molecules. Preferred oligonucleotide primers and cellular sources for amplification therefrom as described in Example 2C.

30 As shown in the Examples, particular combinations of a peptide bound to a recombinant MHC class II molecule used in conjunction with one or more of the above-described recombinant accessory molecules activates T cells into armed effector T

cells that are distinguishable into Th1 inflammatory T cells and Th2 helper T cells.

3. Antigen Processing Assisting Genes and Encoded
5 Molecules

a. HLA-DM

HLA-DM in humans and H2-M in mice is a MHC class II-like molecule that is also encoded within the MHC class II gene clusters. HLA-DM, like MHC class II, contains both α - and
10 β -chain genes forming a heterodimer. However, unlike MHC class II molecules, peptide loading is not required for stabilizing the molecule. HLA-DM facilitates the loading of peptides onto newly formed MHC class II heterodimers following the removal of the invariant chain as further described below. Recombinant
15 HLA-DM is contemplated for use in the compositions and methods of this invention to assist in the loading of internally processed peptides.

b. Invariant Chain

20 The invariant chain is a specialized protein that binds to newly formed MHC class II heterodimers thereby forming a trimer with each subunit of the MHC class II heterodimer. The trimerized molecule prevents the loading of intracellular peptides present in the endoplasmic reticulum but it also
25 facilitates the export of the molecule from that compartment. Thereafter, the invariant chain is cleaved through multiple steps resulting in a MHC class II heterodimer that can then be complexed with processed peptides.

Thus, recombinant invariant chain is contemplated for use
30 in the compositions and methods of this invention to assist in the loading of internally processed peptides.

C. Nucleic Acids and Polynucleotides

1. PCR to Obtain Genes Encoding MHC Class II and
Accessory Molecules

Nucleic acid sequences encoding MHC class II
5 molecules, accessory molecules and antigen processing assisting
molecules of this invention are obtained in a number of ways
familiar to one of ordinary skill in the art including direct
synthesis, cloning, purification of DNA from cells containing
10 such genes, and the like. One expedient means to obtain genes
for encoding the molecules used in the compositions and methods
described herein is by polymerase chain reaction (PCR)
amplification on selected nucleic acid templates with selected
oligonucleotide primer pairs as further described below.

Known, partial and putative human leukocyte antigen (HLA),
15 the genetic designation for the human MHC, amino acid and
nucleotide sequences, including the consensus sequence, are
published (see, e.g., Zemmour and Parham, Immunogenetics 33:
310-320 (1991)), and cell lines expressing HLA variants are
known and generally available as well, many from the American
20 Type Culture Collection ("ATCC"). Therefore, using PCR, MHC
class II-encoding nucleotide sequences are readily operatively
linked to an expression vector of this invention that is then
used to transform an appropriate cell for expression therein.

Particularly preferred methods for producing the
25 recombinant molecules of the present invention rely on the use
of preselected oligonucleotides as primers in PCR to form PCR
reaction products as described herein.

If a gene is to be obtained by PCR amplification, in
general, two primers comprising a PCR primer pair, are used for
30 each strand of nucleic acid to be amplified. For the sake of
simplicity, synthesis of exemplary MHC class II heterodimer-
encoding genes is discussed, but it is expressly to be
understood that the PCR amplification method described is

equally applicable to the synthesis of MHC class II allelic variants, accessory molecules and antigen processing assisting molecules, including those whose complete sequences are presently unknown.

5 In general, a first primer is referred to as a forward primer or a 5' primer as it has the same sequence as the top strand of template DNA and thus hybridizes to the bottom complementary strand.

10 A second primer is referred to as a backward primer or a 3' primer as it has the same sequence as the bottom strand and thus hybridizes to the complementary sequence on the top strand. Typically, in other words, one primer is complementary to the negative (-) or bottom strand of the nucleotide sequence and the other is complementary to the positive (+) or top strand.

15 In preferred aspects, both first and second primers are chosen to hybridize to (i.e., be complementary to) conserved regions within the MHC class II genes. However, primers can be designed to amplify specific MHC class II genes and allelic variants thereof by hybridizing to unique rather than consensus
20 sequences. For this aspect, the template sequence is preferably known for design of such primer pairs.

25 One or both of the first and second primers can be designed to introduce into the amplified product a nucleotide sequence defining an endonuclease recognition site. The site can be heterologous to the MHC class II gene being amplified and typically appears at or near the 5' end of the primer. It may also be helpful to place a 4-base spacer sequence proximal to the restriction site to improve the efficiency of cutting amplification products with enzymes.

30 The primers of the invention for isolating specific nucleotide sequences include oligonucleotides of sufficient length and appropriate sequence so as to provide specific initiation of polymerization on a significant number of nucleic

acids with the corresponding nucleotide sequence. Specifically, the term oligonucleotide primer as used herein refers to a sequence comprising two or more deoxyribonucleotides or ribonucleotides, preferably more than three, and more preferably
5 around 20, the sequence of which is capable of initiating synthesis of a primer extension product.

Experimental conditions conducive to synthesis include the presence of nucleoside triphosphates and an agent for polymerization and extension, such as thermostable polymerases,
10 and a suitable buffer, temperature and pH. The primer is preferably single stranded for maximum efficiency in amplification, but may be double stranded. If double stranded, the primer is first treated to separate the two strands before being used to prepare extension products. Preferably, the
15 primer is an oligodeoxyribonucleotide. The primer must be sufficiently long and substantially complementary to prime the synthesis of extension products in the presence of the inducing agent for polymerization and extension of the nucleotides. The exact length of primer will depend on many factors, including
20 temperature, buffer, and nucleotide composition. The oligonucleotide primer typically contains 15-22 or more nucleotides, although it may contain fewer nucleotides. Alternatively, as is well known in the art, the mixture of nucleoside triphosphates can be biased to influence the
25 formation of mutations to obtain a library of mutated recombinant MHC class II-encoding molecules for use in presenting unique peptides to CD4⁺ T cells.

Oligonucleotide primers of the invention are employed in the PCR amplification process which is an enzymatic chain
30 reaction that produces exponentially growing quantities of a nucleotide sequence. Annealing the primers to denatured nucleic acids followed by extension with a thermostable polymerase such as Thermophilus aquaticus (Taq) and Pyrococcus furiosus (Pfu)

(Hoffman La-Roche, Basal, Switzerland), and nucleotides, results in newly synthesized (+) and (-) strands. Because these newly synthesized sequences are also templates, repeated cycles of denaturing, primer annealing, and extension results in exponential production of the DNA fragment defined by the primers. The product of the chain reaction is a discrete nucleic acid duplex with termini corresponding to the ends of the specific primers employed. Those of skill in the art will know of other amplification methodologies which can also be utilized to increase the copy number of target nucleic acids. These may include for example, ligation activated transcription (LAT), ligase chain reaction (LCR), and strand displacement activation (SDA), although PCR is the preferred method as described in the US Patents listed below.

The oligonucleotide primers of the invention may be prepared using any suitable method, such as conventional phosphotriester and phosphodiester methods as described above for synthesis of complementary oligonucleotides or automated embodiments thereof. One method for synthesizing oligonucleotides on a modified solid support is described in US Patent 4,458,066.

Preferred primers for amplifying MHC class II genes, accessory molecule genes, and antigen processing assisting molecule genes are described in Example 2.

PCR amplification methods are described in detail in U.S. Patent Nos. 4,683,192, 4,683,202, 4,800,159, 4,965,188 and 5,395,750, the disclosures of which are hereby incorporated by reference, and at least in several texts including "PCR Technology: Principles and Applications for DNA Amplification", H. Erlich, ed., Stockton Press, New York (1989); and "PCR Protocols: A Guide to Methods and Applications", Innis et al., eds., Academic Press, San Diego, California (1990). Various preferred methods and primers used herein are described

hereinafter and are also described by Zemmour, et al.,
Immunogenetics, 33:310-20 (1991), by Ausebel, et al., In Current
Protocols in Molecular Biology, Wiley and Sons, New York (1993)
and by Sambrook et al., Molecular Cloning: A Laboratory Manual,
5 Cold Spring Harbor Laboratory, (1989). Particular PCR methods
including nested PCR, overlap PCR, reverse-transcriptase-PCR,
and the like that are well known to one of ordinary skill in the
art are contemplated for use in obtaining the recombinant
molecules of this invention.

10 In alternative embodiments, the PCR process is used not
only to produce a variety of MHC class II-encoding molecules,
but also to induce mutations which may emulate those observed in
the highly-polymorphic MHC loci, or to create diversity from a
single parental clone and thereby provide a MHC class II-
15 encoding DNA "library" having a greater heterogeneity.

2. Expression Vectors

The present invention contemplates plasmid expression
vectors in substantially pure form capable of directing
20 expression of MHC class II-encoding genes, accessory molecule
genes and antigen processing assisting genes to produce the
corresponding recombinant proteins. For simplicity, the above
genes are herein referred to collectively as polypeptide-
encoding nucleotide sequences. Vectors capable of directing the
25 expression of genes to which they are operatively linked are
referred to herein as "expression vectors" or "expression
plasmids", both of which are also referred to as "plasmids".

As used herein, the term "vector" or "plasmid" refers to a
nucleic acid molecule capable of transporting between different
30 genetic environments another nucleic acid to which it has been
operatively linked. Preferred vectors are those capable of
autonomous replication and expression of structural gene
products present in the DNA segments to which they are

operatively linked. Vectors, therefore, preferably contain the replicons and selectable markers that are compatible with the host selection system. One type of preferred vector is an episome, i.e., a nucleic acid capable of extra-chromosomal
5 replication.

A plasmid of this invention is a circular double-stranded plasmid that contains at least a regulation region having elements capable of activating transcription of the translatable polypeptide-encoding nucleotide sequences of this invention.
10 The plasmid further contains a translatable nucleotide sequence from which the desired encoded polypeptides of this invention are expressed. Thus, the vectors are said to be capable of directing the expression of the recombinant polypeptides described herein as encoded from the corresponding expressible
15 genes.

A preferred vector for use according to the present invention is a plasmid; more preferably, it is a high copy number plasmid. It is also preferable that the vector of choice be best suited for expression in the chosen host.

20 Such expression vectors contain a promotor sequence in the regulatory region which facilitates the efficient transcription of an inserted genetic sequence in the host. Preferably, the vector contain an inducible promoter sequence, as inducible promoters tend to limit selection pressure against cells into
25 which such vectors (which are often constructed to carry non-native or chimeric nucleotide sequences) have been introduced. The expression vector also typically contains an origin of replication as well as specific genes which allow phenotypic selection of the transformed cells. The DNA segment
30 can be present in the vector operatively (also referred to as operably) linked to regulatory elements, for example, a promoter (e.g., T7, metallothionein I, or polyhedrin promoters).

In a separate embodiment, a plasmid also contains a gene, the expression of which confers a selective advantage, such as a drug resistance, to a host cell when introduced or transformed into that cell. Typical prokaryotic and eukaryotic drug
5 resistance genes respectively confer resistance to ampicillin or tetracyclin and to neomycin (G418 or Geneticin). Other drug resistance markers include chloramphenicol, kanamycin, streptomycin, carbenicillin, mercury, rifampicin, rifampicin, fusaric acid, and the like.

10 The choice of vector to which the regulatory region and nucleotide sequences for encoding polypeptides of the present invention is operatively linked depends directly, as is well known in the art, on the functional properties desired, e.g., replication or protein expression, and the host cell to be
15 transformed, these being limitations inherent in the art of constructing recombinant DNA molecules.

Operatively linking refers to the covalent joining of nucleotide sequences, preferably by conventional phosphodiester bonds, into one strand of DNA, whether in single or double
20 stranded form. Moreover, the joining of nucleotide sequences results in the joining of functional elements such as response elements in regulatory regions with promoters and downstream polypeptide-encoding nucleotide sequences as described herein.

One typical method for operatively linking inserts into
25 expression plasmids is by directional ligation. This is accomplished through a sequence of nucleotides that are adapted for directional ligation. Such a sequence is referred to commonly as a polylinker that is a region of the DNA expression vector that (1) operatively links for replication and transport
30 the upstream and downstream translatable DNA sequences and (2) provides a site or means for directional ligation of a DNA sequence into the vector. Typically, a directional polylinker is a sequence of nucleotides that defines two or more

restriction endonuclease recognition sequences, or restriction sites. Upon restriction cleavage, the two sites yield cohesive termini to which a translatable DNA sequence can be ligated to the DNA expression vector. Preferably, the two restriction
5 sites provide, upon restriction cleavage, cohesive termini that are non-complementary and thereby permit directional insertion of a translatable DNA sequence into the cassette.

A variety of host-expression vector systems may be utilized to express a polypeptide encoded by a nucleotide sequence.

10 These include but are not limited to microorganisms such as bacteria transformed with recombinant bacteriophage DNA, plasmid DNA or cosmid DNA expression vectors containing a polypeptide-encoding nucleotide sequence; yeast transformed with recombinant yeast expression vectors containing a polypeptide-encoding
15 nucleotide sequence; plant cell systems infected with recombinant virus expression vectors (e.g., cauliflower mosaic virus, CaMV; tobacco mosaic virus, TMV) or transformed with recombinant plasmid expression vectors (e.g., Ti plasmid) containing a polypeptide-encoding nucleotide sequence; insect
20 cell systems infected with recombinant virus expression vectors (e.g., baculovirus) containing a polypeptide-encoding nucleotide sequence; or animal cell systems infected with recombinant virus expression vectors (e.g., retroviruses, adenovirus, vaccinia virus) containing a polypeptide-encoding nucleotide sequence, or
25 transformed animal cell systems engineered for stable expression. In such cases where glycosylation may be important, expression systems that provide for translational and post-translational modifications may be used; e.g., mammalian, insect, yeast or plant expression systems.

30 Any of these systems are useful in practicing the methods of this invention. With any of the above expression systems, the selected host is then used for expression of at least one MHC class II heterodimer alone or in conjunction with at least

one accessory molecule further with or without an antigen processing accessory molecule depending on the actual mechanism for loading peptides, i.e., internally or externally.

Depending on the host/vector system utilized, any of a number of suitable transcription and translation elements, including constitutive and inducible promoters, transcription enhancer elements, transcription terminators, etc. may be used in the expression vector (see e.g., Bitter, et al., Methods in Enzymology, 153:516-544, (1987)). For example, when cloning in bacterial systems, inducible promoters such as P_l of bacteriophage λ , P_{lac}, P_{trp}, P_{tac} (P_{trp-lac} hybrid promoter) and the like may be used. When cloning in mammalian cell systems, promoters derived from the genome of mammalian cells (e.g., metallothionein promoter) or from mammalian viruses (e.g., the retrovirus long terminal repeat; the adenovirus late promoter; the vaccinia virus 7.5K promoter) may be used. Promoters produced by recombinant DNA or synthetic techniques may also be used to provide for transcription of a polypeptide-encoding nucleotide sequence.

In bacterial systems a number of expression vectors may be advantageously selected for expression of a polypeptide-encoding nucleotide sequence according to the methods of this invention. For example, when large quantities are to be produced, vectors which direct the expression of high levels of fusion protein products that are readily purified may be desirable. Those which are engineered to contain a cleavage site to aid in recovering the protein are preferred. Such vectors include but are not limited to the E. coli expression vector pUR278 (Ruther, et al., EMBO J., 2:1791, (1983)), in which the polypeptide-encoding nucleotide sequence may be ligated into the vector in frame with the LacZ coding region so that a hybrid polypeptide-LacZ protein is produced; pIN vectors (Inouye &

Inouye, Nuc. Acids Res., 13:3101-3109, (1985); Van Heeke & Schuster, J. Biol. Chem., 264:5503-5509, (1989)); and the like.

In one embodiment, the vector utilized includes prokaryotic sequences that facilitate the propagation of the vector in
5 bacteria, i.e., a DNA sequence having the ability to direct autonomous replication and maintenance of the recombinant DNA molecule extra-chromosomally when introduced into a bacterial host cell. Such replicons are well known in the art.

Those vectors that include a prokaryotic replicon also
10 typically include convenient restriction sites for insertion of a recombinant DNA molecule of the present invention. Typical of such vector plasmids are pUC8, pUC9, pBR322, and pBR329 available from BioRad Laboratories, (Richmond, CA) and pPL available from Pharmacia, (Piscataway, NJ), and pBLUESCRIPT and
15 pBS available from Stratagene, (La Jolla, CA). A vector of the present invention may also be a Lambda phage vector including those Lambda vectors described in Molecular Cloning: A Laboratory Manual, Second Edition, Maniatis et al., eds., Cold Spring Harbor, NY (1989).

20 In another preferred embodiment, plasmid vectors for use in the present invention are also compatible with eukaryotic cells. Eukaryotic cell expression vectors are well known in the art and are available from several commercial sources. Typically, such vectors provide convenient restriction sites for insertion of
25 the desired recombinant DNA molecule, and further contain promoters for expression of the encoded genes which are capable of expression in the eukaryotic cell, as discussed earlier. Typical of such vectors are pSVO and pKSV-10 (Pharmacia), and pPVV-1/PML2d (International Biotechnology, Inc.), and pTDT1
30 (ATCC, No. 31255).

In addition, in eukaryotic plasmids, one or more transcription units are present that are expressed only in eukaryotic cells. The eukaryotic transcription unit consists of

noncoding sequences and sequences encoding selectable markers. The expression vectors of this invention also contain distinct sequence elements that are required for accurate and efficient polyadenylation. In addition, splicing signals for generating
5 mature mRNA are included in the vector. The eukaryotic plasmid expression vectors can contain viral replicons, the presence of which provides for the increase in the level of expression of cloned genes. A preferred replication sequence is provided by the simian virus 40 or SV40 papovavirus.

10 A preferred expression system for producing the recombinant molecules for use in this invention is an insect system. In one such system, *Autographa californica* nuclear polyhedrosis virus (AcNPV) is used as a vector to express foreign genes. The virus grows in *Spodoptera frugiperda* (Sf9)
15 cells. The polypeptide-encoding nucleotide sequences of this invention may be cloned into non-essential regions (in *Spodoptera frugiperda* for example the polyhedrin gene) of the virus and placed under control of an AcNPV promoter (for example the polyhedrin promoter). Successful insertion of the
20 polypeptide-encoding nucleotide sequence will result in inactivation of the polyhedrin gene and production of non-occluded recombinant virus (i.e., virus lacking the proteinaceous coat coded for by the polyhedrin gene). These recombinant viruses are then used to infect cells in which the
25 inserted gene is expressed. See Smith, et al., J. Biol. Chem., 46:584, (1983); Smith, U.S. Patent No. 4,215,051.

In preferred embodiments, the host cell population is a *Drosophila* cell culture that requires a compatible vector including vectors functionally equivalent to those such as
30 p25-lacZ (see Bello and Couble, Nature, 346:480 (1990)) or pRmHa-1, -2, or -3 (see Bunch, et al., Nucl. Acids Res., 16:1043-1061 (1988)). In the preferred embodiment, the vector is pRmHa-3, which is shown in Figure 1C. This vector includes a

metallothionein promoter, which is preferably upstream of the site at which the MHC sequence is inserted, and the polyadenylation site is preferably downstream of said MHC sequence. Insect cells and, in particular, Drosophila cells are preferred hosts according to the present invention. Drosophila cells such as Schneider-2 (S2) cells, as further described in Section D, have the necessary trans-acting factors required for the activation of the promoter and are thus even more preferred.

The expression vector pRmHa-3 is based on the bacterial plasmid pRmHa-1 (Figure 1A), the latter of which is based on plasmid pUC18 and is deposited with the American Type Culture Collection (ATCC, Rockville, MD), having the accession number 37253. The pRmHa-3 vector contains the promoter, the 5' untranslated leader sequence of the metallothionein gene with the Eco RI and Stu I sites removed as shown in Figure 1C. It also contains the 3' portion of the Drosophila ADH gene including the polyadenylation site. Therefore, cloned DNA is transcriptionally regulated by the metallothionein promoter and polyadenylated. Construction of the pRmHa-1 plasmid is described in Bunch, et al., Nucl. Acids Res. 16: 1043-1061 (1988). Construction of the pRmHa-3 and pRmHa-2 plasmids (the latter of which has a metallothionein promoter sequence that may be removed as an Eco RI fragment) is described in the Examples. With regard to pRmHa-3, a preferred plasmid for use according to the present invention, Pst I, Sph I and Hind III are in the promoter fragment and therefore are not unique. Xba I is in the ADH fragment (4 bases from its 3' end) and is also not unique. The following restriction sites are, however, unique in pRmHa-3 to facilitate cloning of the recombinant genes of this invention: Eco RI, Sac I, Kpn I, Sma I, Bam HI, Sal I, Hinc 2, and Acc I.

Mammalian cell systems that utilize recombinant viruses or viral elements to direct expression may be engineered. For

example, when using adenovirus expression vectors, the coding sequence of a polypeptide may be ligated to an adenovirus transcription/translation control complex, e.g., the late promoter and tripartite leader sequence. This chimeric gene may then be inserted into the adenovirus genome by in vitro or in vivo recombination. Insertion in a non-essential region of the viral genome (e.g., region E1 or E3) will result in a recombinant virus that is viable and capable of expressing the polypeptide in infected hosts (e.g., see Logan & Shenk, Proc. Natl. Acad. Sci., USA, 81:3655-3659, (1984)). Alternatively, the vaccinia virus 7.5K promoter may be used. (e.g., see, Mackett, et al., Proc. Natl. Acad. Sci., USA, 79:7415-7419, (1982); Mackett, et al., J. Virol., 49:857-864, (1984); Panicali, et al., Proc. Natl. Acad. Sci., USA, 79:4927-4931, (1982)). Of particular interest are vectors based on bovine papilloma virus which have the ability to replicate as extrachromosomal elements (Sarver, et al., Mol. Cell. Biol., 1:486, (1981)). Shortly after entry of this DNA into mouse cells, the plasmid replicates to about 100 to 200 copies per cell. Transcription of the inserted cDNA does not require integration of the plasmid into the host's chromosome, thereby yielding a high level of expression. These vectors can be used for stable expression by including a selectable marker in the plasmid, such as the neo gene. Alternatively, the retroviral genome can be modified for use as a vector capable of introducing and directing the expression of a polypeptide-encoding nucleotide sequence of this invention in host cells (Cone & Mulligan, Proc. Natl. Acad. Sci., USA, 81:6349-6353, (1984)). High level expression may also be achieved using inducible promoters, including, but not limited to, the metallothionine IIA promoter and heat shock promoters.

For long-term, high-yield production of recombinant proteins, stable expression is preferred. Rather than using

expression vectors which contain viral origins of replication, host cells can be transformed with a cDNA controlled by appropriate expression control elements (e.g., promoter and enhancer sequences, transcription terminators, polyadenylation sites, etc.), and a selectable marker. As mentioned above, the selectable marker in the recombinant plasmid confers resistance to the selection and allows cells to stably integrate the plasmid into their chromosomes and grow to form foci which in turn can be cloned and expanded into cell lines.

For example, following the introduction of foreign DNA, engineered cells may be allowed to grow for 1-2 days in an enriched media, and then are switched to a selective media. A number of selection systems may be used, including but not limited to the herpes simplex virus thymidine kinase (Wigler, et al., Cell, 11:223, (1977)), hypoxanthine-guanine phosphoribosyltransferase (Szybalska & Szybalski, Proc. Natl. Acad. Sci., USA, 48:2026, (1962)), and adenine phosphoribosyltransferase (Lowy, et al., Cell, 22:817, (1980)) genes, which can be employed in tk⁻, hgp^rt⁻ or ap^rt⁻ cells respectively. Also, antimetabolite resistance-conferring genes can be used as the basis of selection; for example, the genes for dhfr, which confers resistance to methotrexate (Wigler, et al., Proc. Natl. Acad. Sci., USA, 77:3567, (1980); O'Hare, et al., Proc. Natl. Acad. Sci., USA, 78:1527, (1981)); gpt, which confers resistance to mycophenolic acid (Mulligan & Berg, Proc. Natl. Acad. Sci., USA, 78:2072, (1981)); neo, which confers resistance to the aminoglycoside G-418 (Colberre-Garapin, et al., J. Mol. Biol., 150:1, (1981)); and hyg^r, which confers resistance to hygromycin (Santerre, et al., Gene, 30:147, (1984)). Recently, additional selectable genes have been described, namely trpB, which allows cells to utilize indole in place of tryptophan; hisD, which allows cells to utilize histinol in place of histidine (Hartman & Mulligan, Proc. Natl.

Acad. Sci., USA, 85:804, (1988)); and ODC (ornithine decarboxylase) which confers resistance to the ornithine decarboxylase inhibitor, 2-(difluoromethyl)-DL-ornithine, DFMO (McConlogue L., In: Current Communications in Molecular Biology, Cold Spring Harbor Laboratory ed., (1987).

Both prokaryotic and eukaryotic expression vectors are familiar to one of ordinary skill in the art of vector construction and are described by Ausebel, et al., In Current Protocols in Molecular Biology, Wiley and Sons, New York (1993) and by Sambrook et al., Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory, (1989).

For producing recombinant MHC class II α - and β -chains, accessory molecules and antigen processing assisting molecules for use in the compositions and methods of this invention, the respective nucleotide regions are operatively inserted into an expression vector of this invention as described herein. As described in Section B, nucleic acids encoding the recombinant polypeptides of this invention are obtained in a number of ways one of which is by PCR amplification. One of the nucleotide segments to be operatively linked to vector sequences encodes at least a portion of MHC class II α - and β -chains. Preferably, the respective nucleotide sequences for encoding the complete α - and β -chains are separately inserted into an expression vector for expression therefrom; however, it is also feasible to construct a vector which also includes some non-coding MHC sequences as well. The sequences for encoding accessory molecules and antigen processing assisting molecules are similarly inserted into separate expression vectors.

Alternatively, the invention contemplates the presence of more than one polypeptide-encoding gene being present within the same vector, the expression of which is driven by separate regulatory elements, such as a promoter. In other words, the nucleic acids for encoding both α - and β -chains may be

operatively ligated to the same expression vector with or without one or more nucleic acid sequences encoding accessory molecules. Thus, all possible combinations of expression vector construction for producing the recombinant proteins of this invention are contemplated.

In addition to the complete encoding nucleotide sequences as described above, soluble forms of the expressed recombinant polypeptides of this invention are contemplated. The soluble form differs from the non-soluble form in that it contains a "stop" codon inserted prior to the transmembrane domain or other functional location to generate soluble non-membrane anchorable proteins.

D. Synthetic Antigen Presenting Cells and Matrices for Peptide Presentation

1. Synthetic Antigen Presenting Cells and Matrices

In accordance with the present invention, the recombinant MHC class II heterodimers and at least one accessory molecule are operably linked to a matrix comprising a support such that the MHC class II and accessory molecules are present in sufficient numbers to activate a population of CD4⁺ T cells lymphocytes when presented with a peptide complexed to the extracellular portion of the MHC molecule. The peptide can be bound to the MHC class II heterodimer before or after it is linked to the support.

The support can take on many different forms. It can be a solid support such as a plastic or metal material, it can be a porous material such as commonly used in separation columns, it can be a liposome or red blood cell, or it can even be a cell or cell fragment. As discussed in more detail below, in the case where a cell serves as a support, the MHC class II and accessory molecules can be produced by the cell for presentation on that

cell or for presentation on another support that can include a separate cell.

In the former situation, the MHC molecule is then linked to the cell by at least the transmembrane domain if not also the cytoplasmic domain which would not be present in a soluble form of MHC class II. In the latter situation, the extracellular portions of MHC class II molecule and accessory molecule can be linked to a support by providing an epitope which reacts to an antibody immobilized on the support. In addition, the MHC or assisting molecules can be produced with or linked to (His)₆ which would react with nickel in forming part of the support. Other means to immobilize or link MHC molecules to a support are well known in the art.

As discussed above, the support can be a cell membrane or an entire cell. In such a case, an eukaryotic cell line is modified to become a synthetic antigen presenting cell line for use in presenting peptide in the context of MHC class II to T cell lymphocytes. Because empty MHC molecules are thermolabile, it is preferred that the cell culture be poikilotherm and various cell lines are discussed in detail below.

A preferred cell line of the present invention is capable of continuous growth in culture and capable of expressing mammalian MHC class II molecules and accessory molecules on the cell surface. Any of a variety of transformed and non-transformed cells or cell lines are appropriate for this purpose, including bacterial, yeast, insect, and mammalian cell lines. (See, e.g., Current Protocols in Molecular Biology, John Wiley & Sons, NY (1991), for summaries and procedures for culturing and using a variety of cell lines, e.g., E. coli and Sarcomyces cerevisiae).

Preferably, the cell line is a eukaryotic cell line. More preferably, the cell line is poikilothermic (i.e., less sensitive to temperature challenge than mammalian cell lines).

More preferably, it is an insect cell line. Various insect cell lines are available for use according to the present invention, including moth (ATCC CCL 80), armyworm (ATCC CRL 1711), mosquito larvae (ATCC lines CCL 125, CCL 126, CRL 1660, CRL 1591, CRL 6585, CRL 6586), silkworm (ATCC CRL 8851) and butterfly (Spodoptera frugiperda (Sf9 cells, ATCC CRL 1711)). In a preferred embodiment, the cell line is a Drosophila cell line such as a Schneider cell line (see Schneider, J. Embryol. Exp. Morph., 27:353-365 (1972)); preferably, the cell line is a Schneider 2 (S2) cell line (S2/M3) adapted for growth in M3 medium (see Lindquist, et al., Drosophila Information Service, 58:163 (1982)). Schneider 2 (S2) cells have been deposited pursuant to Budapest Treaty requirements with the American Type Culture Collection (ATCC), Rockville, MD, on February 18, 1992, and was assigned accession number CRL 10974.

To generate a synthetic antigen presenting cell of this invention, one or more expression vectors for directing the expression of a selected MHC class II heterodimer in conjunction with one or more accessory molecules is introduced into a recipient host cell. In addition, in alternative embodiments, vectors for expressing antigen processing assisting molecules including HLA-DM and invariant chain are also introduced into the recipient cells. The genes for the above have been described in Sections B and C.

Thus, in order to prepare synthetic antigen presenting cells or matrices, the expression vectors for encoding recombinant polypeptides of this invention are transfected, i.e., introduced, into a selected host cell. The selection of expression vectors as well as the construction thereof is dependent upon the desired outcome for CD4⁺ activation as previously discussed as reiterated below. Transfection, also referred to as transformation, may be accomplished via numerous methods, including the calcium phosphate method, the

DEAE-dextran method, the stable transfer method, electroporation, or via the liposome mediation method. Numerous texts are available which set forth known transfection methods and other procedures for introducing nucleotides into cells; 5 see, e.g., Current Protocols in Molecular Biology, John Wiley & Sons, NY (1991). Following introduction of one or more vectors, the recipient cell is said to be transformed, the selection of which can either be transient or stable.

A culture of cells is first established. A cell line is 10 chosen for transfection because it lacks at least one of the genes being introduced. It has been found that insect cells are advantageous not only because they are poikilothermic, but because they lack these genes and the mechanisms which would otherwise produce MHC molecules bound to peptides. This allows 15 for greater control over the production of peptide-bound MHC molecules, and the production of empty MHC molecules.

The selected cells are then transformed by the introduction of an expression vector containing an expressible MHC class II α -chain gene operably linked to a first promoter and an 20 expressible MHC class II β -chain gene operably linked to a second promoter. A first expressible accessory molecule gene operably linked to a third promoter in a vector is also introduced into the above cell. In a further embodiment, an expressible antigen processing assisting gene operably linked to 25 a fourth promoter in a vector is introduced into the above cell.

In a more preferred embodiment, the vector comprises Drosophila expression plasmid pRMHa-3, described in Section C, into which expressible nucleotide sequences encoding the above 30 recombinant proteins have been inserted using techniques disclosed herein. Preferably, the nucleotide sequences encoding the MHC class II chains, those encoding at least one accessory molecule and those encoding antigen processing assisting

molecules are operably linked to separate expression plasmids that are individually cotransfected into the cultured cells. Alternatively, the nucleotide sequences may be operably linked to separate promoters in the same expression plasmid and
5 cotransfected via that same plasmid. The MHC class II α - and β -chains are preferably from a different species, more preferably, a homeotherm such as mammals and, optimally, humans.

It is preferred that at least one of the genes and, in particular, the MHC class II chain genes be linked to an
10 inducible promoter. This allows control over the production of MHC molecules so that they are only produced at a time when the peptide of interest is available either internally or externally and presented in the culture to react with the produced MHC molecules. This minimizes undesirable MHC molecule/peptide
15 complexes.

Thus, the preferred cell line is a poikilotherm cell line that has separate vectors each containing a MHC class II α - and β -chain gene respectively operably linked to a first and second promoter. Preferably, the promoters are inducible to control
20 the expression of the MHC class II chains. In addition, the cell contains a third vector containing at least a first expressible accessory molecule gene operably lined to a third promoter. In a further embodiment, the cell also contains a fourth vector containing an expressible antigen processing
25 assisting gene operably linked to a fourth promoter. It is preferred that the cell assembles empty MHC molecules and presents them on the cell surface so that the peptides specific for a particular MHC class II haplotype or for a variant allele can be selected as desired.

30 The selection of compatible MHC class II α - and β -chain genes for use in conjunction with one or more particular accessory molecule-encoding genes is dependent upon the T cell activation profile desired. For example, as described in the

Examples, recombinant B7.1 or B7.2 alone or together in conjunction with recombinant murine IA^d MHC class II expressed on the surface of Drosophila APC resulted in proliferation of CD4⁺ T cells having a Th2 profile of increased production of IL-4 and IL-10. In contrast, when either B7.1 or B7.2 were expressed on the surface of Drosophila APC with ICAM-1 along with the same MHC molecules, the activation of CD4⁺ T cells resulted in a Th1 profile with increased IL-2 production and decreased IL-4 and IL-10 production.

Thus, the invention contemplates the production of synthetic APC having on the cell surface any combination of a MHC class II haplotype heterodimer with any one of the accessory molecules of this invention. Particularly preferred combinations include MHC class II with either a costimulatory molecule including B7.1 or B7.2, an adhesion molecule including ICAM-1, ICAM-2, ICAM-3 or LFA-3, or a survival molecule including Fas ligand (FasL). As described above, more than one of each category of accessory molecules can be co-expressed, as for example, B7.1 and B7.2. Alternative preferred embodiments include the permutations where two accessory molecules of different categories are co-expressed on the cell surface. In other words, an adhesion molecule with a costimulatory molecule, and adhesion molecule with a survival molecule, a costimulatory molecule with a survival molecule. In a further embodiment, three accessory molecules of the different categories described herein are co-expressed on the APC surface. It is also contemplated that in all of these embodiments, more than one member of a particular category may be expressed in conjunction with more than one member of other categories. The particular selected combinations are effective on the surface of the cells from which they are expressed or when anchored on the surface of a matrix of this invention. The actual combinations prepared are selected on the basis of T cell activation outcome in view

of the expressed MHC class II molecules complexed with antigenic peptide. Thus, depending on the complex of MHC class II/peptide, the T cell activation outcome may be distinct despite having the same expressed accessory molecules.

5 In a further embodiment, antigen processing assisting genes are co-transfected with any of the above-described combinations to provide for enhanced internal peptide processing and loading. This aspect thus does not involve the generation of empty MHC class II molecules on the cell surface for subsequent peptide
10 complexation. Rather, the expression of invariant chain, HLA-DM or lysosomal enzymes is utilized to allow for optimal processing and loading of proteolytic peptide fragments following cell internalization. The APC of this invention thus can function in either motif of having the recombinant MHC class II heterodimers
15 being loaded either intracellularly or extracellularly.

Successfully transformed cells, i.e., cells that contain at least one expression vector capable of directing the expression of nucleotide sequences according to the present invention, can be identified via well-known techniques. For example, cells
20 resulting from the introduction of a cDNA or rDNA of the present invention can be cloned to produce individual colonies. Cells from those colonies can be harvested, lysed, and their DNA content examined for the presence of the rDNA using a method such as that described by Southern, J. Mol. Biol. 98: 503
25 (1975). In addition to directly assaying for the presence of rDNA, successful transformation or transfection may be confirmed by well-known immunological methods when the rDNA is capable of directing the expression of a subject MHC class II protein of accessory molecule. For example, cells successfully transformed
30 with one or more expression vectors may produce proteins displaying particular antigenic properties which are easily determined using the appropriate antibodies, such as anti-class II of particular haplotypes. In addition, successful

transformation/transfection may be ascertained via the use of an additional vector bearing a marker sequence, such as neomycin resistance, as described hereinabove.

It is also preferable that the culture be stable and
5 capable of sustained growth at reduced temperatures. For example, it is preferred that the culture be maintained at about room temperature, e.g., about 24-27°C. In other embodiments, the culture is maintained at higher temperatures, particularly during the process of activating CD4⁺ cells. It is thus
10 preferred that a culture according to the present invention be capable of withstanding a temperature challenge of about 30°C to about 37°C.

In order to prepare the culture for expression of empty or MHC class II molecules in conjunction with a least one accessory
15 molecule of this invention and optionally antigen processing assisting molecules, the culture may first require stimulation, e.g., via CuSO₄ induction, for a predetermined period of time. After a suitable induction period, e.g., about 12-48 hours, peptides may be added at a predetermined concentration (e.g.,
20 about 0.2 µg/ml to 20 µg/ml). Proteins and peptides for both internal and external loading are prepared as discussed below. After a further incubation period, e.g., for about 12 hours at 27°C, the culture is ready for use in the activation of CD4⁺ cells. While this additional incubation period may be shortened
25 or perhaps omitted, the culture tends to become increasingly stable to temperature challenge if it is allowed to incubate for a time prior to addition of resting or naive CD4⁺ cells. For example, cultures according to the present invention to which peptide has been added are capable of expressing significant
30 amounts of peptide-loaded MHC class II molecules even when incubated for extended periods of time at 37°C.

Nutrient media useful in the culturing of transformed host cells are well known in the art and can be obtained from

numerous commercial sources. In embodiments wherein the host cell is mammalian, a "serum-free" medium is preferably used.

The resulting recombinant expressed MHC class II molecules bind to a particular peptide and are present in sufficient
5 numbers with at least one accessory molecule on the surface of the APC to activate a population of T cell lymphocytes against the MHC class II/peptide complex.

Where the cell line already produces one or more of the desired molecules, it is only necessary to transfect the culture
10 with an expressible gene for the gene which is lacking in the cells. For example, if the cells already present the MHC molecules on their surface, it is only necessary to transfect the culture with a vector containing an expressible gene for the accessory molecule.

15 As previously discussed, a protein or peptide can be introduced into the cell culture at the time the cells are producing MHC class II molecules for internal processing. Through methods such as osmotic shock, the peptides can be introduced in the cell and bind to the produced MHC molecules.
20 Alternatively, particularly in the case poikilotherm cell lines, the MHC molecules will be presented empty on the cell surface. The peptide can then be added to the culture and bound to the MHC molecules as desired. For simplicity, while one peptide is described herein, the methods of this invention contemplate the
25 screening of peptide libraries for identifying novel antigenic peptides for use in the therapeutic methods of this invention.

After the cells are produced having a MHC class II heterodimer and at least one accessory molecules on the cell surface, the cells can be lyophilized to generate cell fragments
30 for use in activating a population of CD4⁺ T cell lymphocytes.

Transfected cultures of cells are also used to produce extracellular portions of MHC class II molecules and accessory molecules. The use of extracellular portions in conjunction

with supports such as solid supports has certain advantages of production. Where living cells are used to provide a synthetic antigen presenting cell, at least three genes, two to produce the MHC class II heterodimer and one for the accessory molecule
5 must be introduced to the cell. Often, additional genes such as for antibiotic resistance are also transfected.

Where a solid support system is being used, one cell line is used to produce the extracellular portions of MHC class II molecules while another cell line is used to produce the
10 extracellular portion of an accessory molecule. The MHC molecule portions and the accessory molecule portions are then harvested from their respective cultures. The molecules are then linked to an appropriate support in sufficient numbers to activate a population of T cells. From a production standpoint,
15 two different cultures can be used, but it is also possible to use the same culture, however, requiring that the culture be transfected with the additional gene for expressing the extracellular portion of an accessory molecule.

A further modification of this embodiment is to provide a
20 third culture of cells which is transfected with an expressible second accessory molecule gene. For example, the second culture of cells produces extracellular portions of the costimulatory molecule while the third culture of cells produce an extracellular portion of an adhesion molecule. The adhesion
25 molecule portions are harvested and linked to the support. In preparing the extracellular portions of a MHC class II heterodimer to be linked to a support, soluble molecules are prepared as previously discussed. These molecules generally lack the transmembrane and cytoplasmic domain in the MHC
30 molecule.

2. Peptides

Virtually all cellular proteins in addition to viral antigens are capable of being used to generate relevant peptide fragments that serve as potential MHC class II-specific peptides. The methods and compositions of this invention provide for MHC class II molecules that have an increased capacity to specifically activate CD4⁺ cells.

The peptides of the present invention bind to MHC class II molecules. The binding occurs under biological conditions which can be created in vivo as well as in vitro. The exact nature of the binding of the peptides need not be known for practice of the invention.

Peptides that bind MHC class II molecules are variable in length and their anchor residues lie at various distances from the ends of the peptide. In one aspect, the peptides prepared for loading onto the MHC molecules are of a single species; i.e., that all peptides loaded onto the MHC be identical in size and sequence so as to produce monoantigenic peptide-loaded MHC class II molecules. In alternative embodiments, peptides are heterogenous and may comprise a random library of peptides to allow for selection of unique members that result in desired T cell activation profiles as previously described. The production and screening of random synthetic peptide libraries is familiar to one of ordinary skill in the art and is described in U.S. Patent Nos. 5,556,762, 5,510,240, 5,498,530, 5,432,018, 5,382,513, 5,338,665 and 5,270,170, the disclosures of which are hereby incorporated by reference.

Peptides may be presented to the cells via various means. Preferably, peptides are presented in a manner which allows them to enter an intracellular pool of peptides. For example, peptides may be presented via osmotic loading. Typically, peptides are added to the culture medium. The peptides may be added to the culture in the form of an intact polypeptide or

protein which is subsequently degraded via cellular processes, e.g., via enzymatic degradation. Alternatively, the intact polypeptide or protein may be degraded via some other means such as chemical digestion (e.g. cyanogen bromide) or proteases (e.g. chymotrypsin) prior to its addition to the cell culture. In other embodiments, the peptides are presented in smaller segments which may or may not comprise antigenic amino acid sequences in conjunction with a particular MHC class II haplotype.

Preferably, a sufficient amount of protein(s) or peptide(s) is added to the cell culture or synthetic matrix to allow the MHC class II molecules to bind and subsequently present a large density of the peptide. preferably, with the same kind of peptide attached to each MHC heterodimer, on the surface of the synthetic APC or matrices of this invention.

In another embodiment of the invention, peptides are added to transfected cells of the present invention in order to enhance the thermostability of the MHC molecules expressed by the cells. As noted above, peptides are preferably added to the culture medium. Antigenic peptides that bind to the MHC class II molecules serve to thermostabilize the MHC molecules and also increase the cell surface expression. Cultures with added peptides which bind to the MHC molecules are thus significantly less susceptible to temperature challenge than cultures without added peptide.

E. Methods of Altering CD4⁺ T Cell Responses

1. Th1 and Th2 CD4⁺ T Cell-Mediated Diseases

Inducing a naive T cell into a desired activated T cell type or deviating the effector function of an activated T cell from a Th1 type to a Th2 type and visa versa is one of the aims of the present invention, especially with regard to

therapeutic methods in treating various CD4⁺ T cell-mediated disease conditions.

The differentiation of a proliferating CD4⁺ T cell into either an inflammatory T cell or a helper T cell is dependent on the cytokines produced by infectious agents, principally IL-12 and IL-4, the influence of accessory molecules and on the nature of the MHC class II/peptide complex. As previously discussed, cell-mediated immunity involves the destruction of intracellular pathogens by macrophages activated by Th1 inflammatory cells directed primarily to intracellular parasites including such parasites as Mycobacterium, Leishmania, Pneumocystis and the like. In contrast, humoral immunity depends on the production of antibody by B cells activated by helper T cells directed primarily at extracellular pathogens including Clostridium, Staphylococcus, Streptococcus, Polio virus, Pneumocystis and the like.

For example, recovery from certain types of infections, such as Leishmania, is associated with preferential production of IL-2/IFN- γ . Mice that mount a Th2 response to Leishmania fail to contain the infection and ultimately die. Inappropriate production of cytokines of the Th2 type response has been frequently linked to allergic type diseases such as asthma and contact sensitivity.

Perhaps the strongest association of human disease with skewed patterns of cytokine production is the association of Th1 responses and Th1 type cytokines with autoimmune disease. Strong evidence in experimental models indicates that many types of autoimmunity including diabetes, experimental models for multiple sclerosis, autoimmune thyroiditis, and the like are mediated by Th1 type CD4⁺ T cells. The expression of Th2-associated cytokines, such as IL-4, in these models interfere with the development of autoimmune disease. Th2 type cytokines

dampen the response of Th1 type cells while the Th1 type cytokines antagonize the development of Th2 type responses.

In view of the association of particular activated T cell subsets with particular disease conditions, a need therefore exists to be able to direct the proliferation and activation of CD4⁺ T cells to a desired T cell subset, a process that is extremely beneficial in altering the course of disease. One potential solution is to activate in vitro CD4⁺ T cells that are first isolated from a subject who may optionally be having either allergy or autoimmune conditions to produce cells secreting a preferred cytokine profile. The resultant activated T cells are then reintroduced to the subject to alter the course of disease and perhaps even provide a long term cure.

Alternative embodiments are directed at the ability to "vaccinate" a potentially responsive individual against the development of either a Th1 or Th2 response, whichever is applicable to that individual. In other words, the selective induction of a particular T cell subset may be achieved by inhibiting naive cells from developing toward the undersired phenotype. For example, in a potentially atopic individual, preventing a deleterious Th2 response would be beneficial as described by Hetzel and Lamb, Clinical Immunol. Immunopath., 73:1-10 (1994).

In still a further embodiment, the compositions and methods of this invention are useful for actively stimulating the development of naive T cells toward the desired phenotype. Existing therapeutic models to date include the use of anti-cytokine antibodies or carrier proteins, the use of idiotypic/GM-CSF fusion protein vaccines to prolong effects of exogenous cytokines, use of selected adjuvants, use of liposome-encapsulated allergens, use of peptide analogs and the like as reviewed by Hetzel and Lamb, id. The authors, however, do state that for chronic Th2 responses to allergens in vivo, little

experimental data exists for the possibility of effecting a desired downregulation of the Th2 response.

In view of the foregoing, the compositions and methods of this invention provide a valuable means to accomplish the
5 therapeutic interventions discussed above.

The present invention allows one to define activation conditions that reproducibly generate CD4⁺ T cell subsets that produce the desired therapeutic cytokine profile. Expression of particular cytokines is linked to a particular antigen presenting cell
10 (APC) and their associated accessory molecules. Since both the cytokines produced by the APC and the coordinately expressed accessory molecules are themselves regulated by multiple factors, including the type of antigen, the affinity of the T cell receptor (TCR)-antigen interaction, antigen concentration
15 and the like, predicting the outcome of T cell activation upon antigen presentation is historically very difficult. Indeed, as additional accessory molecules have been proposed for the activation process in vivo, it has become increasingly clear that many diverse molecules are involved in the regulation of T
20 cell responses and act in combinatorial fashion to effect the outcome of T cell activation.

The present invention provides the generation of synthetic APC that present, in a neutral background, MHC class II molecules in combination with defined accessory molecules that
25 are expressed preferably in a nonmammalian insect cell. The advantage of using the insect cells as the expression and presentation vehicles for the MHC class II/accessory molecule compositions of this invention is that the cells do not endogenously produce regulatory cytokines and do not express
30 mammalian accessory molecules. This overcomes the inherent unpredictability of using mammalian APC that express many molecules that are capable of altering the T cell response. The present invention thus provides the ability to isolate

individual presenting molecules and accessory molecules for expression in selected combinations that permits reproducibility and predictability not available in other approaches.

5 2. Therapeutic Methods

As discussed above, the present invention relates to a method for activating CD4⁺ T cells into differentiated armed effector T cell subtypes. The method relates to providing a synthetic APC or matrix having anchored on the external surface
10 a recombinant MHC class II heterodimer that is capable of binding a peptide. The compositions also have at least one accessory molecule presented on the cell surface. Naive or activated CD4⁺ T cells can be obtained by removal from an individual to be treated. The antigen presenting cells are then
15 contacted with the CD4⁺ T cells for a sufficient period of time to activate the T cells into proliferating and differentiating into a desired T cell phenotype.

The activated CD4⁺ T-cells are separated from the cell line and put into a suspension in an acceptable carrier and
20 administered to the individual.

It is preferred that human genes are used and, therefore, human molecule analogs are produced. As shown in prior U.S. Patent No. 5,314,813, murine systems provide particularly useful models for testing the operation of T cell activation and
25 demonstrate the applicability of the process for human systems. See also Sykulev et al., Immunity, 1:15-22 (1994).

a. Isolation of Resting or Activated CD4⁺ T Cells

Resting (or naive) as well as activated CD4⁺ cells
30 that have not been activated to target a specific antigen presented in the context of MHC class II are extracted from an individual for incubation or exposure to the transformed cultures of the present invention. Naive cells can be

distinguished from primer cells primarily based on the cell surface markers CD455RA and CD45.

It is also preferred that CD4⁺ cells are obtained from an individual prior to the initiation of other treatment or therapy which may interfere with the CD4⁺ cells' ability to be specifically activated. For example, if one is intending to treat an individual with an autoimmune disease, it is preferable to obtain a sample of cells and culture prior to the initiation of adjunctive therapy such as steroid treatment or during a window of time when the patient is not being treated at all.

When activating CD4⁺ T cells to alter the T cell-mediated immune response in a patient, the patient is first analyzed for a patient-specific profile to assess the T cell phenotypic disease state for instituting appropriate counter therapy involving production of the opposing T cell phenotype and cytokines. Cytokine profiles are established with anti-cytokine antibodies that are available from ATCC and by methods described in U.S. Patent Nos. 5,405,751, 5,322,787 and 5,209,920, the disclosures of which are hereby incorporated by reference. Preferred cytokine analyses include interleukin-2 (IL-2), interferon- γ (IFN- γ), tumor necrosis factor (TNF), interleukin-4 (IL-4), interleukin-10 (IL-10) and the like. As previously discussed, particular cytokine profiles are associated with T cell phenotypes and disease states.

In particular, where the condition is an autoimmune disease including multiple sclerosis, autoimmune thyroiditis, systemic lupus erythromatosus, myasthenia gravis, Crohn's disease and inflammatory bowel disease, the cytokine profile is produced by a Th1 type response characterized by increased IL-2, IFN- γ and TNF. In contrast, where the condition is an allergy, such as asthma and contact sensitivity, the cytokine profile is produced by a Th2 type response characterized by increased IL-4 and IL-10.

After analyzing the patient cytokine profile and disease state, patient-isolated CD4⁺ T cells are contacted in vitro with the synthetic APC, cell fragments or matrices of this invention as described below in a sufficient amount for a sufficient time to induce the contacted cells to proliferate and differentiate into activated CD4⁺ T cells that produce a functionally opposing cytokine profile. That is, if the patient were characterized as being a Th1 type responder, the antigen to be presented to the patients CD4⁺ T cells would be that necessary to induce the cells to proliferate and differentiate into a Th2 type. The opposite treatment modality is performed if the patient is characterized with a Th2 type response. Thus, once the opposing activated cells are returned to the patient as described below, the therapeutic goal of effecting an alteration in T cell phenotype response is attained.

Methods of extracting and culturing lymphocytes are well known. For example, U.S. Patent No. 4,690,915 to Rosenberg describes a method of obtaining large numbers of lymphocytes via lymphocytapheresis. Appropriate culturing conditions used are for mammalian cells, which are typically carried out at 37°C.

Various methods are also available for separating out and/or enriching cultures of CD4⁺ cells. Some examples of general methods for cell separation include indirect binding of cells to specifically-coated surfaces. In another example, human peripheral blood lymphocytes (PBL), which include CD4⁺ cells, are isolated by Ficoll-Hypaque gradient centrifugation (Pharmacia, Piscataway, NJ). PBL lymphoblasts may be used immediately thereafter or may be stored in liquid nitrogen after freezing in FBS containing 10% DMSO (Sigma Chemical Co., St. Louis, MO), which conserves cell viability and lymphocyte functions.

Alternative methods of separating out and/or enriching cultures of precursor cells include both positive and negative

selection procedures. For positive selection, after lymphocyte-enriched PBL populations are prepared from whole blood, subpopulations of CD4⁺ lymphocytes are isolated therefrom by affinity-based separation techniques directed at the presence
5 of the CD4 co-receptor antigen. These affinity-based techniques fluorescence-activated cell sorting (FACS), cell adhesion, magnetic bead separation and like methods. (See, e.g., Scher and Mage, in Fundamental Immunology, W.E. Paul, ed., pp. 767-780, River Press, NY (1984).) Affinity methods may utilize
10 anti-CD4 co-receptor antibodies as the source of affinity reagent. Alternatively, the natural ligand, or ligand analogs, of CD4 receptor may be used as the affinity reagent. Various anti-T cell and anti-CD4 monoclonal antibodies for use in these methods are generally available from a variety of commercial
15 sources, including the American Type Culture Collection (Rockville, MD) and Pharmingen (San Diego, CA).

Negative selection procedures are utilized to effect the removal of non-CD4 from the CD4⁺ population. This technique results in the enrichment of CD4⁺ cells from the T and B cell
20 population of leucophoresed patients. Depending upon the antigen designation, different antibodies may be appropriate. For example, monoclonal antibodies OKT4 (anti-CD4, ATCC No. CRL 8002) OKT 5 (ATCC Nos. CRL 8013 and 8016), OKT 8 (anti-CD8, ATCC No. CRL 8014), and OKT 9 (ATCC No. CRL 8021) are identified in
25 the ATCC Catalogue of Cell Lines and Hybridomas (ATCC, Rockville, MD) as being reactive with human T lymphocytes, human T cell subsets, and activated T cells, respectively. Various other antibodies are also available for identifying and isolating T cell species, including precursors and naive and
30 activated memory mature peripheral T cells.

b. In Vitro Activation of CD4⁺ Cells

In order to optimize the in vitro conditions for the generation of specific CD4⁺ T cell phenotypes, the culture of antigen presenting cells is maintained in an appropriate medium.

5 Preferably, when using a support of this invention that is an intact cell, the antigen-presenting cells are Drosophila cells, which are preferably maintained in serum-free medium (e.g. Excell 400). In alternative embodiments, however, when the support is a cell fragment or a matrix of an artificial support
10 as previously described, the culture medium is selected to maintain the viability of the target cells.

Prior to incubation of the synthetic APC, cell fragments or matrices of this invention with the T cells to be activated, an amount of antigenic peptide is provided to the APC or matrices
15 in sufficient quantity to become loaded onto the human MHC class II molecules for expression on the surface of the APC or matrices. As previously discussed, peptide loading can occur intracellularly or extracellularly. Both aspects are accordingly encompassed in the activation process described
20 herein but for simplicity, loading of peptides is generically described as the details of peptide presentation to MHC class II heterodimers and loading have been previously discussed. Moreover, individual peptides as well as peptide libraries are contemplated for use in preparing activated CD4⁺ T cells also as
25 previously described. According to the present invention, a sufficient amount of peptide is an amount that will allow about 200 to about 500,000 and preferably about 200 to 1,000 or more, MHC class II molecules loaded with peptide to be expressed on the surface of each synthetic APC or matrix. Preferably, the
30 above compositions are incubated with 0.2 µg/ml up to 20 µg/ml peptide.

The isolated CD4⁺ cells are then incubated in culture with the appropriate peptide-loaded MHC class II heterodimers

expressed on synthetic APCs or matrices for a time period sufficient to activate CD4⁺ cells. Preferably, the CD4⁺ cells shall thus be activated in an antigen-specific manner. The ratio of CD4⁺ cells to antigen-presenting cells may vary from individual to individual and may further depend upon variables such as the amenability of an individual's lymphocytes to culturing conditions and the nature and severity of the disease condition or other condition for which the within-described treatment modality is used. Preferably, however, the lymphocyte:antigen-presenting cell or matrix ratio is preferably in the range of about 1:1 to 300:1.

The effector/antigen-presenting culture may be maintained for as long a time as is necessary to activate and enrich for a population of a therapeutically useable or effective number of CD4⁺ cells. In general terms, the optimum time is between about one and five days, with a maximum specific level generally being observed after three to five days of culture. In one embodiment of the present invention, in vitro activation of CD4⁺ cells is detected within a brief period of time after transfection of a cell line.

Preferably, the activation of CD4⁺ cells is optimal within one week of exposure to antigen-presenting cells. Thereafter, in a preferred embodiment, the activated CD4⁺ cells are further purified by isolation procedures including density gradients, rosetting with antibody-red blood cell preparations, column chromatography and the like. Following the purification, the resulting CD4⁺ cell preparation is further expanded by maintenance in culture for a period of time to obtain a population of 10⁹ activated CD4⁺ cells. This period may vary depending on the replication time of the cells but may generally be 14 days.

c. Separation of CD4⁺ Cells from Synthetic APC or Matrices

Activated CD4⁺ cells may be effectively separated from the antigen presenting compositions of this invention using one of a variety of known methods. For example, monoclonal antibodies specific for the MHC class II heterodimers, for the peptides loaded thereon, or for the CD4⁺ cells (or a segment thereof) may be utilized to bind their appropriate complementary ligand. Antibody-tagged cells may then be extracted from the stimulator-effector cell admixture via appropriate means such as via well-known flow cytometric or magnetic bead separation methods. Density gradients can also be used to separate blast cells.

d. Therapeutic Treatment with Activated CD4⁺ Cells

Effective amounts of the activated CD4⁺ cells can vary between in vitro and in vivo uses, as well as with the amount and type of target cells expressing the antigenic peptide used in activating the CD4⁺ population. The amount will also vary depending on the condition of the patient and should be determined via consideration of all appropriate factors by the practitioner. Preferably, however, about 1×10^6 to about 1×10^{12} , more preferably about 1×10^8 to about 1×10^{11} , and even more preferably, about 1×10^9 to about 1×10^{10} activated CD4⁺ cells are utilized for adult humans, compared to about 5×10^6 - 5×10^7 cells used in mice.

Preferably, as discussed above, the activated CD4⁺ cells are harvested from the synthetic APC or matrix in culture prior to administration of the CD4⁺ cells to the individual being treated. It is important to note, however, that unlike other present and proposed treatment modalities, in a preferred embodiment, the present method uses a cell culture system of Drosophila APC or acellular matrices that are not tumorigenic. Therefore, if

complete separation of cells and activated CD4⁺ cells is not achieved, there is no inherent danger known to be associated with the administration of a small number of synthetic APC or matrices, whereas administration of mammalian tumor-promoting
5 cells may be extremely hazardous.

Methods of re-introducing cellular components are known in the art and include procedures such as those exemplified in U.S. Patent No. 4,844,893 to Honsik, et al. and U.S. Patent No. 4,690,915 to Rosenberg. For example, administration of
10 activated CD4⁺ cells via intravenous infusion is appropriate.

Examples

The following examples are intended to illustrate, but not limit, the present invention.

15

1. Preparation of pRmHa-3 Expression Vector

The pRmHa-3 expression vector for use in expressing MHC proteins in Drosophila Schneider 2 (S2) cells as described in
20 this invention was constructed by ligating a Sph I-linearized pRmHa-1 DNA expression vector with a DNA fragment resulting from a Sph I restriction digest of a pRmHa-2 expression vector to form the pRmHa-3 expression vector as described below. The ligating of the linearized pRmHa-1 with the pRmHa-2 fragment in
25 this manner was performed to remove one of two Eco RI restriction endonuclease cloning sites present in pRmHa-1. Thus, the resultant pRmHa-3 expression vector contained only one Eco RI restriction site in the multiple cloning site (polylinker) into which various MHC class II-encoding DNA
30 fragments were inserted as described in the Examples.

A. Preparation of pRmHa-1 Expression Vector

The pRmHa-1 expression vector, containing a metallothionein promoter, metal response consensus sequences (designated MT) and an alcohol dehydrogenase (ADH) gene containing a polyadenylation signal isolated from Drosophila melanogaster, was constructed as described by Bunch et al., Nucl. Acids Res., 16:1043-61 (1988). The plasmid expression vector, pUC18, having the ATCC accession number 37253, was used as the source vector from which subsequent vectors described herein were derived. The pUC18 plasmid contains the following restriction sites from 5' to 3' in the multiple cloning site, all of which are not illustrated in the schematic representations of the pUC18-derived vectors in Figure 1: Eco RI; Sac I; Kpn I; Sma I and Sma I located at the same position; Bam HI; Xba I; Sal I, Acc I and Hinc II located at the same position; Pst I; Sph I and Hind III. The pUC18 vector was first digested with Hind III to form a linearized pUC18. Blunt ends were then created by filling in the Hind III ends with DNA polymerase I large fragment as described by Maniatis et al., Molecular Cloning: A Laboratory Manual, eds. Cold Spring Harbor Laboratory, New York (1982).

The resultant linearized blunt-ended pUC18 vector was ligated with a 740 base pair (bp) Hinf I fragment from the Drosophila melanogaster ADH gene containing a polyadenylation signal. The ligated ADH allele was first isolated from the plasmid pSACI, described by Goldberg et al., Proc. Natl. Acad. Sci., USA, 77:5794-5798 (1980), by digestion with Hinf I followed by blunt ending with Klenow resulting in the nucleotide sequence listed in SEQ ID NO 1. The pSACI vector containing the ADH allele was constructed by subcloning into pBR322 (ATCC accession number 31344) a 4.7 kilobase (kb) Eco RI fragment of Drosophila DNA selected from a bacteriophage lambda library containing random, high molecular weight (greater than 15 kb).

The 5' Hinf I restriction site occurred naturally in the ADH gene at position 1770 as described by Kreitman, Nature, 304:412-417 (1983). The 3' Hinf I site was derived from the pUC18 vector into which the ADH gene had been cloned. This position was four bases 3' to the Xba I site at position 2500 of the ADH gene. The ADH segment extended from 35 bp upstream of the polyadenylation/cleavage sequence in the 3' untranslated portion of the ADH mRNA to 700 bp downstream of the polyadenylation signal. The resultant pUC18-derived vector containing the ADH gene fragment was designated pHa-1 as shown in Figure 1A.

A 421 bp Eco RI/Stu I MT gene fragment for insertion into pHa-1 was obtained from a clone containing DNA of approximately 15.3 kb in a Drosophila melanogaster genomic DNA library. The library, prepared with a Mbo I partial digestion of imaginal DNA, was cloned in the lambda derivative EMBL4. The 421 bp fragment contained the MT promoter and metal response consensus elements of the Drosophila MT gene (Maroni et al., Genetics, 112:493-504 (1986)). This region, containing the promoter and transcription start site at nucleotide position +1, corresponded to position -370 to nucleotide position +54 of the MT gene (SEQ ID NO 2). The resultant fragment was then ligated into pHa-1 expression vector prepared above that was previously linearized with Eco RI and Sma I. The 3' blunt end in MT created by the Stu I digest was compatible with the blunt end in pHa-1 created by the Sma I digest. The resultant pUC18-derived vector containing a 5' Drosophila MT gene fragment and a 3' ADH gene fragment was designated pRmHa-1. The pRmHa-1 expression vector contained the origin of replication (ori) and the beta-lactamase gene conferring resistance to ampicillin (Amp^r) from pUC18 as shown in Figure 1A on the pHa-1 vector. The pRmHa-1 also contained from 5' to 3' the MT gene fragment, the multiple cloning site and the ADH gene fragment. The pRmHa-1 vector was

used as described below in the construction of the pRmHa-3 expression vector.

B. Preparation of pRmHa-2 Expression Vector

5 For constructing the pRmHa-2 expression vector shown in Figure 1A, the MT fragment prepared above was inserted into the pUC18-derived vector pHa-1 as described for constructing pRmHa-1 above with a few modifications. An Eco RI linker was added to the Stu I site of the Eco RI/Stu I-isolated MT gene
10 fragment prepared above to form a metallothionein fragment having Eco RI restriction sites on both ends. The resultant fragment was then ligated into the ADH fragment-containing pUC18 expression vector that was previously linearized with Eco RI. The resultant pUC18-derived vector containing a 5' Drosophila MT
15 gene fragment and a 3' ADH gene fragment having two Eco RI restriction sites 5' to the multiple cloning site was designated pRmHa-2. The pRmHa-2 expression vector contained the origin of replication (ori) and the beta-lactamase gene conferring resistance to ampicillin (Amp^r) from pUC18. The diagram of
20 pRmHa-2 also shows the 5' to 3' contiguous positions of the MT gene fragment, the multiple cloning site and the ADH gene fragment. The pRmHa-2 vector was used along with pRmHa-1 as described below in the construction of the pRmHa-3 expression vector.

25

C. Preparation of pRmHa-3 Expression Vector

To prepare the pRmHa-3 expression vector that had only one Eco RI restriction site, a fragment from pRmHa-2 was ligated into pRmHa-1. For this construction, pRmHa-2, prepared above,
30 was first digested with Sph I. The resultant Sph I fragment beginning in the middle of the MT gene and extending to the Sph I site in the multiple cloning site was first isolated from the pRmHa-2 vector and then ligated into pRmHa-1 that was previously

modified to remove the Eco RI restriction site 5' to the MT gene fragment then linearized with Sph I. This process is schematically illustrated in Figure 1B. To remove the Eco RI site in pRmHa-1, the vector was first digested with Eco RI to
5 form a linearized vector, then blunt ended with Mung Bean nuclease and religated.

A schematic of the pRmHa-3 vector is shown in Figure 1C. The relative positions of the various restriction sites from the pUC18 vector from which pRmHa-3 was derived are indicated in the
10 figure. The pRmHa-3 vector, being derived from pUC18, contains the pUC18 origin of replication and beta-lactamase gene conferring ampicillin resistance. Thus, MHC class II-encoding DNA fragments as prepared in this invention and cloned into the multiple cloning site of pRmHa-3 were transcriptionally
15 regulated by the MT promoter and polyadenylated via the ADH gene.

2. Preparation and Expression of Expressible MHC Class II Genes

20 A. Amplification and Expression of MHC Class II Genes

Genes or cDNAs encoding any preferred mammalian MHC Class II are cloned via use of the polymerase chain reaction (PCR). The primers described herein are used to amplify the appropriate Class II cDNAs in separate reactions which are then
25 cloned and sequenced. To create MHC class II proteins, full length murine IA^d α - and β - chain cDNAs were first obtained followed by PCR amplification and modification. The complete nucleotide sequences of the murine IA^d α -chain cDNA is described by Benoist et al., Cell, 34:169-177 (1983) and is also listed in
30 Genebank Accession Number K01923. The complete nucleotide sequences of the murine IA^d β -chain cDNA is described by Malissen et al., Science, 221:750-754 (1983) and is also listed in GenBank Accession Numbers K00007 and K00008. To amplify each

chain, mouse splenocytes were used as a source of total RNA. First strand cDNA was synthesized by using oligo(dT) and avian myeloblastosis virus reverse transcriptase. The resulting cDNA was used in a PCR amplification reaction utilizing the

5 appropriate primers described below and a GeneAmp kit and thermal cycler (Perkin-Elmer/Cetus, Norwalk, CT). Reaction conditions preferably contained 1 μ g cDNA template and 200 nM of each oligonucleotide primer. Thirty cycles were run as follows: (a) 1 minute at 92°C; (b) 1 minute at 60°C; and (c) 1 minute at

10 72°C. The PCR reaction was then heated to 99°C for 10 minutes to inactivate the Taq polymerase and the ends of the DNA were made blunt by T4 polymerase (Stratagene, La Jolla, CA).

For all the oligonucleotide primers indicated in the Examples, the 5' primer is also referred to as the forward

15 primer or sense primer as it has the same sequence as the top strand of the cDNA for hybridizing to the complementary bottom strand. In contrast, the 3' primer is also referred to as the backward primer or anti-sense primer as it has the same sequence as the bottom strand of the cDNA for hybridizing to the

20 complementary top strand. Both primers are written in the 5' to 3' direction. The full length IA^d α -chain cDNA was amplified with the 5' primer having the nucleotide sequence

5'CTTGAATTCCACCATGCCGTGCAGCAGAGCTCTGA3' (SEQ ID NO 3). The 5' primer was also designed with an Eco RI restriction site to

25 allow for directional ligation of the amplified products into recipient expression vectors. The 3' primer

5'TTTGGATCCTCATAAAGGCCCTGGGTGTC3' (SEQ ID NO 4) was also designed to contain a Bam HI restriction site.

The full length IA^d β cDNA was amplified with the 5' forward

30 primer having the nucleotide sequence

5'CTTGAATTCCACCATGGCTCTGCAGATCCCCA3' (SEQ ID NO 5). The 5' primer was also designed with an Eco RI restriction site to allow for directional ligation of the amplified products into

recipient expression vectors. The 3' primer
5'TTTGGATCCTCACTGCAGGAGCCCTGCT3' (SEQ ID NO 6) was designed to
contain a Bam HI restriction site. The modified cDNAs encoding
the murine IA^d α- and β-chains were first separately
5 directionally cloned into the polylinker at the Eco RI and Bam
HI restriction sites of the metallothionein promoter-driven
pRmHa-3 vector, then sequenced by dye terminator technique on an
Applied Biosystem 373A automated sequencer.

The complete nucleotide sequences of the murine IA^d α- and
10 β-chain amplified regions cloned into pRmHa-3 are respectively
listed in SEQ ID NOS 7 and 8.

The separate plasmids were then transfected into Drosophila
melanogaster Schneider-2 cells (ATCC CRL 10974, Rockville, MD)
as described elsewhere (Jackson et al., Proc. Natl. Acad. Sci.
15 U.S.A., 89;12117-21 1992). Equal amounts of the plasmids
encoding the two class II chains were cotransfected together
with a neomycin resistance gene, plasmid phshsneo (Bunch et al.,
Nuc. Acids Res., 16:1043-1061 (1988) at a ratio of 1:30 to
produce stable cell lines that were derived by G418 selection in
20 Schneider's Drosophila medium (Gibco/BRL, Grand Island, NY)
supplemented with 10% fetal calf serum (heat treated for 1 hour
at 55°C), 100 units/ml penicillin, 100 µg/ml streptomycin, and 1
mM glutamine over a period of 4 weeks. Specifically, after
transfection, the supernatant was carefully removed and the
25 cells were transferred to a 75cm² flask in a total volume of 12
ml Schneider medium containing 500 µg/ml Geneticin (G418)
(Gibco/BRL, Grand Island, NY). After 4 days, 4 ml of the
culture were transferred to a fresh flask containing 6 ml of
Schneider medium with 500 µg/ml G418. This procedure was
30 repeated every 4-7 days until a stable population of cells
emerged which weakly adhered to the flask and grew with a
doubling time of approximately 24 hours. These cells were
subsequently cultured and passaged in the selection media as

described above. Frozen aliquots of the stably transfected cells were prepared by collecting 5-20 X 10⁶ cells by centrifugation and resuspending them in 1 ml of cell freezing media (93% fetal calf serum/7% dimethylsulfoxide). Aliquots
5 were then placed at -70°C for 1 week and subsequently transferred to liquid nitrogen storage. Expression of the stably transfected MHC class II IA^d α- and β-chain genes was induced with 0.7 mM cupric sulfate for 24 hours at 27°C.

Expression of MHC class II IA^d heterodimers at the cell
10 surface of the transfected Drosophila cells was evaluated by flow cytometry after staining with with MKD6, an IA^d-specific monoclonal antibody (Kappler et al., J. Exp. Med., 153:1198 (1981). Briefly, aliquots of cells (5 X 10⁵) were transferred into tubes on ice, collected by centrifugation (1,000 X g for 5
15 minutes), resuspended in 0.1 ml of Drosophila medium with 5% horse serum containing the appropriate primary antibody (MKD6). After a 20 minute incubation at room temperature, cells were washed twice in 3 ml of Schneider's medium containing horse serum and resuspended in 0.1 ml of medium containing FITC-
20 labeled secondary antibody (Cappell, Durham, NC). After a 20 minute incubation on ice, cells were washed twice with Schneider's medium containing horse serum and resuspended in this buffer at a concentration of 1 X 10⁶/ml. Propidium iodide was added to permit exclusion of dead cells from the analysis.
25 Samples were then analyzed by FACScan or FACSsort instrument (Becton Dickinson). Synthetic antigen presenting cells of this invention that express the murine IA^d MHC class II haplotype, described herein and well known in the art, with one or more accessory molecules were produced as described below.

30 Full-length human MHC class II α- and β-chains for the DR, DQ and DP haplotypes are amplified from peripheral blood cells which include B cells, macrophages and dendritic cells. The 5'

and 3' primers used for amplifying each of the haplotypes have the following sequences:

DR α : 5' primer = 5'CCACCATGGCCATTAGTGGAGTC3' (SEQ ID NO 9)

3' primer = 5'TTTGGATCCTTACAGAGGCCCCCTGCGTT3' (SEQ ID NO 10);

DR β : 5' primer = 5'CCACCATGGTGTGTCTGAGGCTCC3' (SEQ ID NO 11)

3' primer = 5'TTTGGATCCTCAGCTCAGGAATCCTCTTG3' (SEQ ID NO 12);

DQ α : 5' primer = 5'CCACCATGGTCCTAAACAAAGCTCTGAT3' (SEQ ID NO 13)

3' primer = 5'TTTGGATCCTCACAAGGGCCCTTGGTGTCT3' (SEQ ID NO 14);

DQ β : 5' primer = 5'CCACCATGGCTTGGAAGAAGGCCTTT3' (SEQ ID NO 15)

3' primer = 5'TTTAGATCTCAGTGCAGAAGCCCTTT3' (SEQ ID NO 16);

DP α : 5' primer = 5'CCACCATGGGCCCTGAAGACAGAAT3' (SEQ ID NO 17)

3' primer = 5'TTTGGATCCTCACAGGGTCCCCTGGGC3' (SEQ ID NO 18);

DP β : 5' primer = 5'CCACCATGGTTCTGCAGGTTTCTGCG3' (SEQ ID NO 19)

3' primer = 5'TTTGGATCCTTATGCAGATCCTCGTTGAA3' (SEQ ID NO 20).

The amplification conditions for obtaining the above human haplotypes are identical to those described above for the murine counterparts.

25 Synthetic antigen presenting cells of this invention that express a human MHC class II haplotypes, described herein and well known in the art, with one or more accessory molecules are produced as described below.

30 B. Amplification of Invariant and HLA-DM Chain cDNAs

For some aspects of the present invention as described in Section B in the detailed description, the invariant chain and HLA-DM α and HLA-DM β are co-expressed with a selected MHC

class II heterodimer and at least one accessory molecule described above. In this instance, the MHC class II α - and β -chains, invariant chain and HLA-DM α - and β -chains are transfected into recipient antigen presenting cells in molar ratios 5:5:8:1:1. The ratio of accessory molecules to the molar ratio of the other genes is 1:1. The resultant transfected cells are then induced for expression of the exogenous genes as described above to generate a synthetic antigen presenting cell line of this invention.

10

1) Invariant Chain

The murine invariant chain cDNA is generated from mouse spleen cells. The full-length invariant chain is amplified from this source with the primer pair of a 5' and 3' primer having the respective nucleotide sequences 5'AAGAATTCAGTAGAGGCTAGAGCCAT3' (SEQ ID NO 21) and 5'AAGGATCCTCACAGGGTGACTTGACC3' (SEQ ID NO 22). As described above, the 5' and 3' primers were designed to respectively incorporate Eco RI and Bam HI restriction sites. Following cloning of the amplified invariant chain into a Eco RI/Bam HI-digested pRmHa-3 vector, the resultant clone was sequenced and determined to have the sequence listed in SEQ ID NO 23.

The human invariant chain cDNA used in this invention is generated using RNA from γ -interferon-induced HeLa cells (ATCC Accession Number CCL 2). The resultant cDNA is then used as a template in PCR with the 5' and 3' oligonucleotide primers having the respective nucleotide sequences 5'AAGAATTCACCATGGATGATCAGCGGACCTT3' (SEQ ID NO 24) and 5'AAAGGATCCTCACATGGGGACTGGGCCCAGA3' (SEQ ID NO 25). The resulting PCR fragments are cleaved with Eco RI and Bam HI before ligation into similarly digested pRmHa-3.

2) HLA-DM

Messenger RNA from γ -interferon-induced HeLa cells is also used to synthesize the α - and β -chains of HLA DM cDNA that is then used as a template in PCR. The α -chain is amplified with the 5' and 3' primer pair having the respective nucleotide sequences 5'AAACCATGGGTCATGAACAGAACCA3' (SEQ ID NO 26) and 5'TTGTCGACTCAGTCACCTGAGCAAGG3' (SEQ ID NO 27). The β -chain is amplified with the 5' and 3' primer pair having the respective nucleotide sequences 5'AAACCATGGTCTCATTCCTGCC3' (SEQ ID NO 28) and 5'TTGTCGACCTAGGAAATGTGCCATCC3' (SEQ ID NO 29). The resulting PCR fragments are cleaved with Nco I and Sal I before separate ligation into a similarly digested pRmHa-3.

C. Amplification of Genes Encoding Accessory Molecules1) Adhesion Moleculesa. ICAM-1, ICAM-2, and ICAM-3

For isolating murine ICAM-1, spleen cells were isolated from Balb/c mice. The spleen cells were first stimulated with conA before isolation of mRNA using the FastTrack kit (Invitrogen, San Diego, CA) according to the manufacturers' instructions. cDNA was synthesized from the mRNA as described above. The resultant cDNA was then subjected to PCR with the following respective 5' and 3' primers that were designed based on the published cDNA nucleotide sequence (Siu, G. et al., J. Immunol., 143:3813-3820 (1989)):

5'TTTAGAATTACCATGGCTTCAACCCGTGCCAAG3' (SEQ ID NO 30) and

5'TTTAGTCGACTCAGGGAGGTGGGGCTTGTCC3' (SEQ ID NO 31). The PCR

products were then cleaved with the restriction enzymes Eco RI and Sal I and ligated into a similarly digested pRmHa-3.

The above expression construct was then co-transfected into Drosophila S2 cells with the murine IA^d α - and β -chain genes prepared above using the calcium phosphate method as described above in the molar ratio of 1:1:1 of ICAM-1: α -chain: β -chain.

Stably transfected cells were obtained as described above. Synthetic antigen presenting Drosophila S2 cells expressing the accessory molecule ICAM-1 with the IA^d α- and β-chains on the surface of the cell were then produced by inducing expression as previously described.

In other embodiments, synthetic antigen presenting cells containing genes encoding ICAM-1, B7.1 and/or B7.2 in conjunction with the murine IA^d molecules were also produced for generating synthetic antigen presenting cells as used in activation assays as described in Example 5.

Human ICAM-1 is similarly amplified from mRNA isolated from the human cell line K562, originated from human chronic myelogenous leukemia (ATCC Accession Number CCL-243) and cultured under recommended conditions (i.e., RPMI with 10% fetal calf serum at 37°C with 5% CO₂). The PCR primers for amplifying the human accessory molecules are designed based on known available sequences and in consideration of the 5' and 3' cloning sites needed to clone into the appropriate vectors. The nucleotide sequence of human ICAM-1 cDNA is available through GenBank Accession Number GB J03132. The ICAM-1 5' and 3' primers have the respective nucleotide sequences 5'ACCCTTGAATTCATGGCTCCCAGCAGCCCCCGGCC3' (SEQ ID NO 32) and 5'ATTACCGGATCCTCAGGGAGGCGTGGCTTGTGTGTTCCGG3' (SEQ ID NO 33).

PCR is performed with these primers as previously described to obtain amplified human ICAM-1. The resultant PCR products are then cloned into pRmHa-3 as previously described followed by transfection into recipient cells, also as previously described.

Similarly, human ICAM-2 and ICAM-3, the nucleotide sequences of which are available with the respective GenBank Accession Numbers GB X15606 and GB S50015, are amplified with comparable primer pairs. The respective 5' and 3' primers for amplifying ICAM-2 have the nucleotide sequence

5 'AAGGTACCCGTGGAGACTGCCAGAGAT3' (SEQ ID NO 34) and
5 'TTTGGATCCCTATGGCCGGAAGGCCTG3' (SEQ ID NO 35). The respective
5' and 3' primers for amplifying ICAM-3 have the nucleotide
sequence 5 'AAGAATTCCTGTCAGAATGGCCACCAT3' (SEQ ID NO 36) and
5 5 'TTTAGATCTTCACTCAGCTCTGGACGGT' (SEQ ID NO 37). The resultant
amplified ICAM products are then separately ligated into pRmHa-3
and separately transfected into Drosophila S2 cells.

Synthetic antigen presenting Drosophila S2 cells expressing
the accessory molecule ICAM-1 with a selected human MHC class II
10 haplotype on the surface of the cell are then produced by
inducing expression as previously described.

In other embodiments, synthetic antigen presenting cells
containing genes encoding ICAM-1, ICAM-2 or ICAM-3 in
combination with B7.1 and/or B7.2 along with a human haplotype
15 are also produced for generating synthetic antigen presenting
cells as used in activation assays as described in Example 5.
Additional permutations contemplated for use in this invention
include the above combinations with LFA-3 and/or Fas ligand
(FasL), both of which are other accessory molecules as described
20 below.

b. LFA-3

Human LFA-3 is isolated from blood
lymphocytes. The nucleotide sequence of human LFA-3 cDNA is
available through GenBank Accession Number GB I09083. Human
25 LFA-3 is amplified accordingly with a 5' and 3' primer having
the respective nucleotide sequences
5 'ACCCTTGAGCTCATGGTTGCTGGGAGCGACGCGGG3' (SEQ ID NO 38) and
5 'ATTACCGGATCCTTAAAGAACATTCATATACAGCACAATACA3' (SEQ ID NO 39).

As described previously, synthetic antigen presenting cells
30 containing genes encoding human LFA-3 alone or in various
combinations with the other accessory molecules described herein
along with a human haplotype are also produced for generating

synthetic antigen presenting cells as used in activation assays as described in Example 5.

2) Costimulatory Molecules

a. B7.1

5 cDNA was generated from mRNA isolated from mice as described above for ICAM-1. The resultant cDNA was then subjected to PCR with the following respective 5' and 3' oligonucleotide primers shown in the 5' to 3' direction that were designed based on the published cDNA nucleotide sequence
10 (Freeman, et al., J. Exp. Med., 174:625-631 (1991)):
5'TTTAGAATTCACCATGGCTTGCAATTGTCAGTTG3' (SEQ ID NO 40) and
5'TTTAGTCGACCTAAAGGAAGACGGTCTGTTC3' (SEQ ID NO 41). The PCR products were cleaved with the restriction enzymes Eco RI and Sal I and ligated into a similarly digested pRmHa-3.

15 The above expression construct was then co-transfected into Drosophila S2 cells with the murine IA^d α- and β-chain genes prepared above using the calcium phosphate method as described above in the molar ratio of 1:1:1 of B7.1:α-chain:β-chain. Stably transfected cells were obtained as described above.
20 Synthetic antigen presenting Drosophila S2 cells expressing the accessory molecule B7.1 with the IA^d α- and β-chains on the surface of the cell were then produced by inducing expression as previously described.

In other embodiments, synthetic antigen presenting cells
25 containing genes encoding murine ICAM-1, B7.1 and/or B7.2 in conjunction with the murine IA^d molecules are also produced for generated synthetic antigen presenting cells as used in activation assays as described in Example 5.

Human B7.1 is similarly isolated from K562 cells and cloned
30 into pRmHa-3 as described above. The nucleotide sequence of human B7.1 cDNA is available through GenBank Accession Number GB M83071. The 5' and 3' primers have the respective nucleotide sequences, 5'ACCCTTGAATCCATGGGCCACACACGGAGGCAG3' (SEQ ID NO 42)

and 5'ATTACCGGATCCTTATACAGGGCGTACACTTTCCTTCT3' (SEQ ID NO 43). The resultant PCR products are then inserted into pRmHa-3 that is then co-transfected along with amplified cloned human haplotype genes into Drosophila S2 cells.

- 5 Synthetic antigen presenting Drosophila S2 cells expressing the accessory molecule B7.1 with a selected human MHC class II haplotype on the surface of the cell are then produced by inducing expression as previously described.

In other embodiments, synthetic antigen presenting cells
10 containing genes encoding human B7.1 in various combinations with the other accessory molecules described herein along with a human haplotype are also produced for generating synthetic antigen presenting cells as used in activation assays as described in Example 5.

15

b. B7.2

Murine IC-21 cells (ATCC Accession Number TIB 186) were propagated in RPMI 1640 medium containing 10% fetal calf serum. cDNA was synthesized from the mRNA isolated
20 from these cells as described above. The resultant cDNA was then subjected to PCR with the following respective 5' and 3' oligonucleotide primers shown in the 5' to 3' direction that were designed based on the published cDNA nucleotide sequence (Freeman, et al., J. Exp. Med., 178:2185-2192 (1993)):

- 25 5'TTTAGAATTACCATGGACCCAGATGCACCATGGG3' (SEQ ID NO 44) and 5'TTTAGTCGACTCACTCTGCATTTGGTTTGTCTGA3' (SEQ ID NO 45). The PCR products were cleaved with the restriction enzymes Eco RI and Sal I and ligated into a similarly digested pRmHa-3.

The above expression construct was then co-transfected into
30 Drosophila S2 cells with the murine IA^d α- and β-chain genes prepared above using the calcium phosphate method as described above in the molar ratio of 1:1:1 of B7.2:α-chain:β-chain. Stably transfected cells were obtained as described above.

Synthetic antigen presenting Drosophila S2 cells expressing the accessory molecule B7.2 with the IA^d α - and β -chains on the surface of the cell were then produced by inducing expression as previously described.

5 In other embodiments, synthetic antigen presenting cells containing genes encoding murine B7.2 in conjunction with ICAM-1 and the murine IA^d molecules were also produced following the procedures described above for generated synthetic antigen presenting cells as used in activation assays as described in
10 Example 5.

Human B7.2 is isolated from the human cell line HL60, that originated from a human promyelocytic leukemia (ATCC Accession Number CCL-240). The nucleotide sequence of human B7.2 cDNA is available through GenBank Accession Number GB M83071. The 5'
15 and 3' primers for amplifying human B7.2 have the respective nucleotide sequences 5'ACCCTTGAGCTCATGGATCCCCAGTGCACTATG3' (SEQ ID NO 46) and 5'ATTACCCCCGGGTAAAAACATGTATCACTTTTGTTCGCATGA3' (SEQ ID NO 47).

The amplified B7.2 products are then cloned into pRmHa-3 as
20 previously described for tranfection into Drosophila S2 cells along with constructs for a selected human haplotype. Expression of the transfected genes is induced as previously described.

In other embodiments, synthetic antigen presenting cells
25 containing genes encoding human B7.2 in various combinations with the other accessory molecules described herein along with a human haplotype are also produced for generating synthetic antigen presenting cells as used in activation assays as described in Example 5.

30

3. Survival Molecules

Human Fas ligand is isolated from activated human T cells. The nucleotide sequence of human Fas ligand cDNA is

available through GenBank Accession Number GB U08137. Human Fas ligand is amplified accordingly with a 5' and 3' primer pair having the respective nucleotide sequences

5'AAAGGATCCACCATGCAGCAGCCCTTCAATT3' (SEQ ID NO 48) and

5'TTTGGATCCTTAGAGCTTATATAAGCCGA3' (SEQ ID NO 49).

Human CD70, the ligand for CD27 expressed on T cells, is amplified as described above with a 5' and 3' primer pair having the respective nucleotide sequences

5'AAAGAATTCGGTACCATGCCGGAGGAGGGTTCGG3' (SEQ ID NO 50) and

5'TTTGGATCCTCAGGGGCGCACCCTACTGCA3' (SEQ ID NO 51).

As described previously, synthetic antigen presenting cells containing genes encoding human Fas ligand alone or in various combinations with the other accessory molecules described herein along with a human haplotype are also produced for generating synthetic antigen presenting cells as used in activation assays as described in Example 5.

3. Preparation of MHC Class II Anchored on Synthetic Supports

The examples described herein present a new method to immobilize high amounts of MHC class II molecules and single accessory molecules or various combinations thereof on various surfaces (fly cells, red blood cells, latex beads) in native conformation as judged by monoclonal antibody binding and rosetting experiments (T cell receptor binding). This method can be extended to other synthetic surfaces including artificial phospholipid membranes. Phosphatidylethanolamine as well as avidin-coupled phospholipids are particularly relevant this invention. These phospholipids are commercially available from Lipex Biomembrane Inc., Vancouver, BC, Canada.

A. Immobilization of Biotinylated MHC Class II on Avidin-Coated Red Blood Cells

NHS-LC-biotin, neutravidin and biotin-BMCC are purchased from Pierce (Rockford, IL). Sheep red blood cells are
5 obtained from the Colorado Serum Company (Denver, CO).
Drosophila S2 cells expressing MHC class II are prepared as described above. Monoclonal antibodies MKD6 and M5-114 (ATCC Accession Number TIB 120 for the hybridoma secreting MKD6) are used as hybridoma cell culture supernatants. The protocol used
10 is described by Muzykantov and Taylor (Anal. Biochem., 223:142-148 (1994)). Briefly, SRBC are washed four times in PBS, biotinylated using NHS-LC-biotin, washed again 4 times in PBS, incubated with neutravidin, and finally washed four times and stored at 4°C in PBS containing 3% fetal calf serum and
15 0.02% sodium azide. Recombinant expressed MHC class II α - and β -chains are biotinylated using biotin-BMCC, a maleimide-coupled biotin which reacts with thiol groups. Biotinylation is performed as recommended by the manufacturer. Unreacted biotin is removed using Centricon 10.
20 Biotinylated MHC class II α - and β -chains are immobilized by incubation at a final concentration of 0.2 mg/ml with avidin-coated SRBC for 30 minutes followed by washing in DMEM containing 10% fetal calf serum. SRBC with attached MHC class II are used immediately.
25 Immobilization of biotinylated MHC class II α - and β -chains on avidin-coated SRBC is done as indicated above. Attachment is assessed using flow cytometry using the antibodies described above. For rosetting assays, either Drosophila S2 cells expressing MHC class II or MHC class II-coated SRBC are
30 incubated with specific peptide (see Example 4) (0.02 mM) or an irrelevant peptide (0.02 mM) for 30 minutes on ice; CD4+ T cells are then added, the proportion being 10 T cells for one Drosophila S2 cell, or 10 SRBC for 1 T cell. The mixture is

then pelleted and kept on ice for at least 30 minutes. Cells are then carefully resuspended and rosettes are counted, a rosette being a Drosophila S2 cell bound to at least 3 CD4+ T cells, or a CD4+ T cell bound to at least 3 SRBC.

5

B. Immobilization of Biotinylated MHC Class II on Avidin-Coated Latex Beads

Six micron diameter latex sulfate beads are purchased from Interfacial Dynamics Corporation (Portland, OR) and
10 biotinylated according to the protocol described above. Avidin-coated latex beads are prepared using a 1% suspension of the latex beads incubated in PBS containing 1 mg/ml of neutravidin for 1 hour at room temperature. An equal volume of PBS containing 10% fetal calf serum is then added. After 1 hour
15 of incubation at room temperature, the beads are washed 3 times and used for binding of recombinant biotinylated MHC class II.

Recombinant biotinylated MHC class II is immobilized by incubation at a final concentration of 0.2 mg/ml with avidin-coated latex beads for 30 minutes followed by washing in
20 DMEM containing 10% fetal calf serum. SRBC with attached MHC class II are used immediately. Rosetting experiments are performed as previously described.

C. Immobilization and Detection of MHC Class II Bound to Various Solid Supports Such as Plastic Microwell Plates

The MHC class II molecules are immobilized by direct binding to microtiter plates (Corning) and detected as follows: MHC class II is diluted to a desired concentration in PBS, e.g.
30 1 mg/ml for 100 ng/well, and then added to each well on the plastic microtiter plate. The plate is incubated for 1 hour at room temperature. After incubation, the plate is washed once with PBS and 200 μ l 2% bovine serum albumin (BSA) in PBS +

(.05%) and Tween (PBST) is added followed by incubation for another hour at room temperature. The plate is washed 3 times with PBST and biotinylated anti-MHC class II monoclonal antibody was added (1:2500) in 2% BSA in PBS. The plate is incubated
5 another hour at room temperature and washed 3 times with PBST. Avidin conjugated HRP is added (1:2500) in 2% BSA in PBS. Following another hour of incubation at room temperature, the plate is washed 3 times with PBST and H_2O_2 or thophenyldiamine was added. The reaction is stopped with H_2SO_4 . Reaction product
10 is detected colorimetrically at 490 nm. Recombinant MHC class II molecules can alternatively be bound through biotin-avidin linked interactions with the substrate. In this embodiment, the microwell plates are coated with 100 μ l avidin diluted in PBS to a concentration of 0.001 mg/ml. Excess avidin is removed by a
15 PBS wash. The above procedure for presenting and detecting MHC class II binding is followed.

Recombinant MHC molecules are alternatively immobilized by a linkage based on a poly-histidine tag added to the MHC interacting with the nickel bound to the substrate.

20 The above procedure for binding and detection is followed using nickel chelate coated microwell plates (Xenopore) and expressed recombinant MHC molecules with a poly-histidine tag.

4. Peptide Generation

25 Antigenic peptides according to the present invention are obtained from naturally-occurring sources, including those described in the specification, or may be synthesized using known methods. In various examples disclosed herein, all peptides were separately chemically synthesized by T-BOC
30 chemistry on an Applied Biosystem instrument, ABI 431A (Foster City, CA), and purified on a C18 reverse phase column. Isolation or synthesis of peptides generated from a random assortment of amino acid residues may also be appropriate,

particularly when one is attempting to ascertain a particular epitope in order to load an empty MHC molecule with a peptide most likely to stimulate CD4 cells. Mixtures of randomized peptides are obtained by using proteasomes, by subjecting a
5 protein or polypeptide to a degradative process, such as digestion with chymotrypsin, or by synthesis.

While the cell lines of the present invention are able to degrade proteins and polypeptides into smaller peptides capable of being loaded onto human MHC class II molecules, it is
10 preferable to introduce the peptides directly into the cell culture to facilitate a more rapid loading and expression process.

If peptides are synthesized having randomly incorporated amino acid residues, all varieties of amino acids, L or D
15 conformations as well as modified amino acids, are preferably incorporated during each cycle of the synthesis. It should be noted, however, that various parameters, e.g., solvent incompatibility of certain amino acids, may result in a mixture which contains peptides lacking certain amino acids. The
20 process should thus be adjusted as needed, by altering solvents and reaction conditions, to produce the greatest variety of peptides. A peptide used for peptide loading as described in Example 5 with murine MHC class II is ovalbumin₃₂₃₋₃₃₉ (OVA) having the amino acid residue sequence ISQAVHAAHAEINEAGR (SEQ ID NO
25 52). Preferred peptides specific for human MHC class II haplotypes include the following peptides: 1) Influenza hemagglutinin₃₀₆₋₃₁₈ PKYVKQNTLKLAT (SEQ ID NO 53) that binds HLA-DRB1*0101; 2) HSP65₂₋₁₂ from M. tuberculosis KTITATDEEARR (SEQ ID NO 54) that binds HLA-DRB1*0301; 3) Mitochondrial outer protein
30 (MOMP) of C. trachomatis QASLALSYRLNMFTP (SEQ ID NO 55) that binds HLA-DRB1*0301; and 4) HLA-A2₃₃₋₄₅ FVRFDSAASQRM (SEQ ID NO 56) that binds to HLA-DRB1*0401. An extensive listing of known class II-binding peptides as well as a description of how

peptide ligands are defined is presented by Rammensee et al., Immunogenetics, 41:178-228 (1995). Peptides for use in practicing the methods of this invention are also available through Chiron Mimetopes. In other embodiments, peptide libraries, readily produced by one of ordinary skill in the art, are contemplated for use in practicing the methods of this invention with the synthetic antigen presenting cells as described herein.

10 5. Loading of Empty MHC Class II Molecules by In Vitro Incubation with Peptides

As shown by the results presented below, the murine MHC class II molecules along with selected accessory molecules expressed on the surface of Drosophila cell lines, having been
15 transfected with various combinations of plasmids encoding MHC class II and accessory molecules, bind specific peptides in a dose-dependent fashion and are able to present these peptides to antigen specific T cells resulting in T cell activation.

For the proliferation and activation assays performed
20 below, Drosophila melanogaster Schneider 2 (SC2) cells (ATCC Accession Number CRL 10974) were transformed with genes encoding the MHC class II α - and β -chain from H2-A^d as described in Example 2. For other assays where selected accessory cells were expressed in combination with the MHC class II molecules, the
25 genes for B7.1, B7.2 and ICAM-1, were also introduced into the MHC gene-containing cells as previously described.

The accessory molecules were expressed at selected concentrations and in combinations as defined below to permit reproducible and consistent activation of CD4⁺ cells. Following
30 expression, induced by treatment with copper sulfate, the concentrations of expressed molecules were monitored. Selection of cells expressing the desired levels of molecules was performed with flow cytometry sorting of cell lines. Table I

lists the Drosophila cell lines expressing murine IA^d MHC class II molecules alone or in combination with various accessory molecules. The table also presents a summary of the results of proliferative effects, described further below, as indicated with a "-" for no effect and a "+" for a positive proliferative effect. A summary of the types of cytokines produced by each type of synthetic antigen presenting cell is also indicated, and described further below, with the additional notation with arrows of whether a particular cytokine production was up or down regulated.

Table 1

	<u>Accessory</u> <u>Molecules</u>	<u>Proliferative</u> <u>Responses</u>	<u>Cytokines</u> <u>Produced</u>
15	None	-	-
	B7.1	+	IL-4, IL-10, IL-2
	B7.2	+	IL-4, IL-10, IL-2
	ICAM-1	-	-
	B7.1 + B7.2	+	IL-4, IL-10, IL-2
20	B7.1 + ICAM-1	+	↑IL-2, ↓IL-4, ↓IL-10
	B7.2 + ICAM-1	+	↑IL-2, ↓IL-4, ↓IL-10
	CD70 + ICAM-1	+	IL-2
	CD70	-	-
	CD70 + ICAM-1		
25	+ B7.2	+	↑IL-2

For assessing the effect of antigen presentation and ultimately T cell response, the antigen specific T cells were CD4⁺ cells obtained from T cell receptor transgenic mice. The transgenic line as used herein was D01 which expresses a T cell receptor specific for the ovalbumin peptide 323-339. The D01 mouse line is obtained from Dr. Dennis Lo. For assaying T cell responsiveness to the recombinant surface-expressed MHC class II

molecules, CD4⁺ cells were first purified from lymph nodes using a combination of techniques designed to eliminate all host APC from the responder population. Small numbers (5×10^4) of these highly purified CD4⁺ cells were then cultured with the Drosophila cell lines in the presence or absence of ova peptide.

The effect of peptide presentation in the presence of MHC class II with and without accessory molecules, B7.1, B7.2 or B7.2 plus ICAM-1, was first assessed in a proliferation assay performed as described by Webb et al., Cell, 63:1249 (1990). Briefly, 5×10^4 CD4⁺ T cells were cultured in microtiter wells with 5×10^4 Drosophila cells in 300 μ l of culture medium. Eighteen hours prior to harvesting, 100 μ l of supernatant were removed for cytokine analysis and 1 μ Ci of ³H-thymidine was added to each well. At the times indicated (3, 4 or 5 days), the wells were individually harvested onto glass fiber filters and counted in a liquid scintillation counter.

As seen in Figure 2, cell lines expressing MHC class II molecules alone failed to stimulate T cell proliferative responses as indicated by the low incorporation of radioactive thymidine; higher counts indicate greater proliferation plotted against the amount of stimulating peptide in μ M. However, co-expression of either B7.1 or B7.2 costimulatory molecules conferred the capacity of MHC class II presented-ova peptide to stimulate strong proliferative responses. For comparison, the results with splenic APC (labeled T-S) are shown in Figure 2.

As shown in Figure 3, Drosophila cell lines expressing MHC class II molecules with the adhesion molecule ICAM-1 were nonstimulatory. However, the expression of ICAM-1 together with B7.2 generated a synthetic APC cell line that stimulated proliferative responses at lower concentrations of peptide antigen. These results demonstrate that Drosophila cell lines expressing the murine H2-A^d MHC class II molecules and costimulatory molecules, B7.1, B7.2 and B7.2 plus ICAM-1, are

able to activate naive CD4⁺ T cells as measured by an increase in proliferation.

In addition to proliferation, the reproducible activation of CD4⁺ T cells to produce particular cytokines was desired.

5 Thus, the cytokines produced in the cultures described above were also evaluated using standard ELISA methods with antibodies specific for the cytokine to be measured. The data plotted in Figures 4A-4D and 5A-5D indicate the amount of cytokine produced in ng/ml concentrations against the stimulatory molecules on
10 APC.

The results of these assays indicate that different combinations of accessory molecules on Drosophila APC expressing recombinant MHC class II leads to the production of different cytokines as shown in Figures 4A-4D and Figures 5A-5D. For IL-2
15 production, a Th1 type response, cell lines expressing MHC class II with B7.1 or B7.2 were poorly stimulatory as shown in Figure 4A. In order to achieve significant levels of IL-2, the Drosophila cell lines required co-expression of ICAM-1, B7 and class II (Figure 5A). Production of γ -interferon was also
20 poorly induced by Drosophila cell lines expressing B7.1 and B7.2 (Figure 4D). In contrast to the results with IL-2, however, the co-expression of ICAM-1 did not restore γ -IFN production (Figure 5C). Splenic APC (labeled as T-S) induced strong γ -IFN production suggesting that additional molecules regulate the
25 production of this cytokine.

IL-4 production, a Th2 type response, is generally not observed during primary stimulation of naive CD4⁺ T cells. In keeping with these observations, splenic APC (T-S) failed to induce significant IL-4 after culture with D01 cells and
30 ova-peptide (Figure 5B). However, Drosophila cells co-expressing class II MHC molecules with either B7.1 or B7.2 or both induced strong IL-4 production during culture (Figures 4B and 5B). Interestingly, the addition of ICAM-1 had an

antagonistic influence on the production of this cytokine (Figure 5B). IL-10 was similarly regulated to IL-4 as shown in Figures 4D and 5D.

As described above, the results with a cytokine profile of a Th2 type response, rather than a Th1 type response, were obtained with cell lines expressing B7 and MHC class II. In contrast, co-expression of MHC class II in combination with B7 and ICAM-1 antagonized the production of IL-4 and IL-10 while strongly promoting IL-2, thereby inducing a Th1 type response with support the present invention in the ability to drive the activation of CD4⁺ T cells into desired T cell subsets. These results definitively show that Drosophila cell lines can be tailored to express defined combinations of accessory molecules for use in achieving a desired population of CD4⁺ T cells that produce different patterns of cytokines. Thus, the compositions and methods described herein validate the compositions and methods of use of the present invention.

The foregoing is intended to be illustrative of the present invention, but not limiting. Numerous variations and modifications may be effected without departing from the true spirit and scope of the invention.

What is claimed is:

1. A synthetic antigen presenting cell (APC) for activating CD4⁺ T cells comprising:

a) a MHC class II α -chain gene operably linked to a first promoter in a vector capable of expressing a MHC class II α -chain;

b) a MHC class II β -chain gene operably linked to a second promoter in a vector capable of expressing a MHC class II β -chain, wherein upon expression of the α -chain and β -chain genes, the α -chain and β -chain form a MHC class II heterodimer capable of loading a peptide; and

c) at least one accessory molecule gene operably linked to a third promoter in a vector capable of expressing an accessory molecule, wherein at least one of the Class II genes and accessory molecule gene is lacking from the APC.

2. The APC of claim 1 wherein the α - and β -chain genes are of human origin.

3. The APC of claim 1 wherein at least one promoter is inducible.

4. The APC of claim 1 wherein the α -, β - and accessory molecule genes are present in the same vector.

5. The APC of claim 1 wherein at least one of the α -, β - and accessory molecule genes are present in a separate vector.

6. The APC of claim 1 wherein the APC is an insect cell.

7. The APC of claim 6 wherein the insect cell is selected from the group consisting of Spodoptera and Drosophila.

8. The APC of claim 1 further comprising a neomycin resistance gene operably linked to a vector.

9. The APC of claim 1 wherein the accessory molecule gene encodes a costimulatory molecule.

10. The APC of claim 9 wherein the costimulatory molecule is B7.1 or B7.2.

11. The APC of claim 1 wherein the accessory molecule gene encodes an adhesion molecule.

12. The APC of claim 11 wherein the adhesion molecule is ICAM-1, ICAM-2, ICAM-3, or LFA-3.

13. The APC of claim 1 wherein the accessory molecule gene encodes a survival molecule.

14. The APC of claim 13 wherein the survival molecule is Fas ligand or CD70.

15. The APC of claim 1 having a gene for a first accessory molecule and a gene for a second accessory molecule.

16. The APC of claim 15 wherein the first accessory molecule is a costimulatory molecule and the second accessory molecule is an adhesion molecule.

17. The APC of claim 16 wherein the costimulatory molecule is B7.1 or B7.2 and the adhesion molecule is ICAM-1.

18. The APC of claim 15 wherein the first accessory molecule is a costimulatory molecule and the second accessory molecule is a survival molecule.

19. The APC of claim 15 wherein the first accessory molecule is a survival molecule and the second accessory molecule is an adhesion molecule.

20. The APC of claim 19 wherein the survival molecule is CD70 and the adhesion molecule is ICAM-1.

21. The APC of claim 15 wherein the first and second accessory molecules are costimulatory molecules.

22. The APC of claim 21 wherein the costimulatory molecules are B7.1 and B7.2.

23. The APC of claim 1 having a gene for a first accessory molecule, a gene for a second accessory molecule and a gene for a third accessory molecule.

24. The APC of claim 23 wherein the first accessory molecule is a costimulatory molecule, the second accessory

molecule is an adhesion molecule, and the third accessory molecule is a survival molecule.

25. The APC of claim 24 wherein the costimulatory molecule is B7.2, the adhesion molecule is ICAM-1 and the survival molecule is CD70.

26. The APC of claim 1 wherein the MHC class II heterodimer and accessory molecule are present on the external surface of the APC in sufficient numbers for activating CD4⁺ T cells when a peptide is loaded onto the heterodimer.

27. The APC of claim 26 wherein the peptide is loaded extracellularly.

28. The APC of claim 26 wherein the peptide is loaded intracellularly.

29. The APC of claim 1 further comprising an antigen processing assisting gene operably linked to a fourth promoter in a vector capable of expressing an antigen processing assisting molecule.

30. A cell fragment derived from the APC of claim 1 having the MHC class II heterodimer and at least one accessory molecule operably associated on the fragment for activating CD4⁺ T cells.

31. The cell fragment of claim 30 wherein the MHC class II heterodimer is empty.

32. The cell fragment of claim 30 wherein a peptide is loaded onto the MHC class II heterodimer.

33. A synthetic antigen presenting matrix for activating CD4⁺ T cells comprising:

- a) a support;
- b) an extracellular portion of a MHC class II heterodimer operably linked to the support and capable of loading a selected peptide; and
- c) an extracellular portion of at least one accessory molecule operably linked to the support such that the

extracellular portions of the MHC class II heterodimer and accessory molecule are present on the matrix in sufficient numbers for activating CD4⁺ T cells when a peptide is loaded onto the extracellular portion of the heterodimer.

34. The matrix of claim 33 wherein the support is a cell fragment.

35. The matrix of claim 33 wherein the support is a cell.

36. The matrix of claim 35 wherein the extracellular portion of the MHC molecule is linked to the cell by a transmembrane domain of the MHC class II heterodimer.

37. The matrix of claim 33 wherein the support is a liposome.

38. The matrix of claim 33 wherein the support is a solid surface.

39. The matrix of claim 33 wherein the extracellular portion of the MHC class II heterodimer is linked to an epitope which reacts with an antibody to link the portion to the support.

40. The matrix of claim 33 wherein the extracellular portion of the Class II MHC heterodimer is linked to (His)₆ which reacts with nickel to link the portion to the support.

41. The matrix of claim 33 wherein the support is a porous material.

42. The matrix of claim 33 wherein the peptide is loaded onto the extracellular portion of the MHC class II heterodimer.

43. The matrix of claim 33 wherein the extracellular portion of the MHC class II heterodimer is empty.

44. The matrix of claim 33 wherein the accessory molecule is a costimulatory molecule.

45. The matrix of claim 44 wherein the costimulatory molecule is B7.1 or B7.2.

46. The matrix of claim 33 wherein the accessory molecule is an adhesion molecule.

47. The matrix of claim 46 wherein the adhesion molecule is ICAM-1, ICAM-2, ICAM-3 or LFA-3.

48. The matrix of claim 33 wherein the accessory molecule is a survival molecule.

49. The matrix of claim 48 wherein the survival molecule is Fas ligand or CD70.

50. The matrix of claim 33 having a first accessory molecule and a second accessory molecule.

51. The matrix of claim 50 wherein the first accessory molecule is a costimulatory molecule and the second accessory molecule is an adhesion molecule.

52. The matrix of claim 51 wherein the costimulatory molecule is B7.1 or B7.2 and the adhesion molecule is ICAM-1.

53. The matrix of claim 50 wherein the first accessory molecule is a costimulatory molecule and the second accessory molecule is a survival molecule.

54. The matrix of claim 50 wherein the first accessory molecule is a survival molecule and the second accessory molecule is an adhesion molecule.

55. The matrix of claim 54 wherein the survival molecule is CD70 and the adhesion molecule is ICAM-1.

56. The matrix of claim 50 wherein the first and second accessory molecules are costimulatory molecules.

57. The matrix of claim 56 wherein the costimulatory molecules are B7.1 and B7.2.

58. The matrix of claim 50 further comprising a third accessory molecule.

59. The matrix of claim 58 wherein the first accessory molecule is a costimulatory molecule, the second accessory molecule is an adhesion molecule, and the third accessory molecule is a survival molecule.

60. The matrix of claim 59 wherein the costimulatory molecule is B7.2, the adhesion molecule is ICAM-1 and the survival molecule is CD70.

61. A method of producing a synthetic antigen presenting cell (APC) comprising:

- a) transforming a cell with an expressible MHC class II α -chain gene operably linked to a first promoter in a vector capable of expressing a MHC class II α -chain;
- b) transforming a cell with an expressible MHC class II β -chain gene operably linked to a second promoter in a vector capable of expressing a MHC class II β -chain; and
- c) transforming a cell with a first expressible accessory molecule gene operably linked to a third promoter in a vector capable of expressing an accessory molecule.

62. The method of claim 61 wherein the cell lacks a gene coding for at least one of the α -chain, the β -chain and the accessory molecule genes.

63. The method of claim 61 further comprising the step of transforming the cell with an expressible antigen processing assisting gene operably linked to a fourth promoter in a vector capable of expressing an antigen processing assisting molecule.

64. The method of claim 61 wherein the α - and β - chain genes are of human origin.

65. The method of claim 61 wherein at least one promoter is inducible.

66. The method of claim 61 wherein the α -, β - and accessory molecule genes are present in the same vector.

67. The method of claim 61 wherein the α -, β - and accessory molecule genes are present in separate vectors.

68. The method of claim 61 wherein the cell is an insect cell.

69. The method of claim 68 wherein the insect cell is selected from the group consisting of Spodoptera and Drosophila.

70. The method of claim 61 further comprising the step of transforming the cell with an expressible neomycin resistance gene operably linked to a vector.

71. The method of claim 61 wherein the accessory molecule gene encodes a costimulatory molecule.

72. The method of claim 71 wherein the costimulatory molecule is B7.1 or B7.2.

73. The method of claim 61 wherein the accessory molecule gene encodes an adhesion molecule.

74. The method of claim 73 wherein the adhesion molecule is ICAM-1, ICAM-2, ICAM-3 or LFA-3.

75. The method of claim 61 wherein the accessory molecule gene encodes a survival molecule.

76. The method of claim 75 wherein the survival molecule is Fas ligand or CD70.

77. The method of claim 61 further comprising the step of transforming the cell with a gene for a second accessory molecule.

78. The method of claim 77 wherein the first accessory molecule is a costimulatory molecule and the second accessory molecule is an adhesion molecule.

79. The method of claim 77 wherein the first accessory molecule is a costimulatory molecule and the second accessory molecule is a survival molecule.

80. The method of claim 77 wherein the first accessory molecule is a survival molecule and the second accessory molecule is an adhesion molecule.

81. The method of claim 77 further comprising the step of transforming the cell with a gene for a third accessory molecule.

82. The method of claim 81 wherein the first accessory molecule is a costimulatory molecule, the second accessory

molecule is an adhesion molecule, and the third accessory molecule is a survival molecule.

83. A method of producing a synthetic antigen presenting cell (APC) comprising:

a) providing a cell lacking a gene encoding at least one of MHC class II α -chain, MHC class II β -chain, and an accessory molecule; and

b) transforming the cell with an expressible gene for each of the genes of (a) lacking in the cell, the gene being operably linked to a promoter in a vector capable of expressing the gene.

84. A method of producing a synthetic antigen presenting cell (APC) comprising:

a) providing a cell lacking a gene encoding at least one of MHC class II α -chain, MHC class II β -chain, an accessory molecule and an antigen processing assisting molecule; and

b) transforming the cell with an expressible gene for each of the genes of (a) lacking in the cell, the gene being linked to a first operable promoter in a vector capable of expressing the gene.

85. A method of producing a synthetic antigen matrix comprising:

a) providing an extracellular portion of a recombinant MHC class II heterodimer;

b) providing an extracellular portion of at least one recombinant accessory molecule; and

c) linking the MHC class II heterodimer and accessory molecule to a support in sufficient numbers for activating CD4⁺ T cells when a peptide is loaded onto the MHC class II heterodimer.

86. The method of claim 85 wherein the accessory molecule is a costimulatory molecule.

87. The method of claim 86 wherein the costimulatory molecule is B7.1 or B7.2.

88. The method of claim 86 wherein the accessory molecule is an adhesion molecule.

89. The method of claim 88 wherein the adhesion molecule is ICAM-1, ICAM-2, ICAM-3 or LFA-3.

90. The method of claim 85 wherein the accessory molecule is a survival molecule.

91. The method of claim 90 wherein the survival molecule is Fas ligand or CD70.

92. A method for activating CD4⁺ T cells in vitro, the method comprising:

- a) providing the APC of claim 26;
- b) contacting the APC of step a) with CD4⁺ T cells, thereby inducing the contacted CD4⁺ T cells to proliferate and differentiate into activated CD4⁺ T cells.

93. The method of claim 92 further comprising:

- c) separating the activated CD4⁺ T cells from the APC.

94. The method of claim 93 further comprising the step of adding the activated CD4⁺ T cells to an acceptable carrier or excipient to form a suspension.

95. The method of claim 94 further comprising the step of administering the suspension to a patient.

96. A method for activating CD4⁺ T cells in vitro, the method comprising:

- a) providing the cell fragment of claim 30;
- b) loading the MHC class II heterodimer with a peptide; and
- c) contacting the peptide-loaded cell fragment with the CD4⁺ T cells, thereby inducing the contacted CD4⁺ T cells to proliferate and differentiate into activated CD4⁺ T cells.

97. The method of claim 96 further comprising the step of separating the activated CD4⁺ T cells from the cell fragment.

98. The method of claim 97 further comprising the step of adding the activated CD4⁺ T cells to an acceptable carrier or excipient to form a suspension.

99. The method of claim 98 further comprising the step of administering the suspension to a patient.

100. A method for activating CD4⁺ T cells in vitro, the method comprising:

- a) providing the matrix of claim 33;
- b) loading the MHC class II heterodimer with a peptide; and
- c) contacting the peptide-loaded cell matrix with the CD4⁺ T cells, thereby inducing the contacted CD4⁺ T cells to proliferate and differentiate into activated CD4⁺ T cells.

101. The method of claim 100 further comprising the step of separating the activated CD4⁺ T cells from the matrix.

102. The method of claim 101 further comprising the step of adding the activated CD4⁺ T cells to an acceptable carrier or excipient to form a suspension.

103. The method of claim 102 further comprising the step of administering the suspension to a patient.

104. A method for activating CD4⁺ T cells in vitro, the method comprising:

- a) contacting the APC of claim 1, the cell fragment of claim 30, or the matrix of claim 33, in an amount sufficient with a peptide library in vitro for a sufficient time to generate a peptide-loaded MHC class II heterodimer;
- b) contacting the peptide-loaded MHC class II heterodimer of step b) with CD4⁺ T cells, thereby inducing the contacted CD4⁺ T cells to proliferate and differentiate into activated CD4⁺ T cells.

105. A method of altering a CD4⁺ T cell-mediated immune response to treat a condition in a patient comprising:

- a) analyzing the patient for patient-specific cytokine profile;
- b) collecting CD4⁺ T cells from the patient;
- c) contacting the CD4⁺ T cells with the APC of claim 26 in vitro in a sufficient amount and for a sufficient time, thereby inducing the contacted CD4⁺ T cells to proliferate and differentiate into activated CD4⁺ T cells that produce a functionally opposing cytokine profile to the profile obtained in step a); and
- d) returning the activated CD4⁺ T-cells to the patient.

106. The method of claim 105 wherein the condition is an autoimmune disease.

107. The method of claim 106 wherein the autoimmune disease is selected from the group consisting of diabetes, multiple sclerosis, autoimmune thyroiditis, systemic lupus erythromatosus, mysasthenia gravis, Crohn's disease and inflammatory bowel disease.

108. The method of claim 106 wherein the patient-specific cytokine profile is produced by a CD4⁺ Th1 type response.

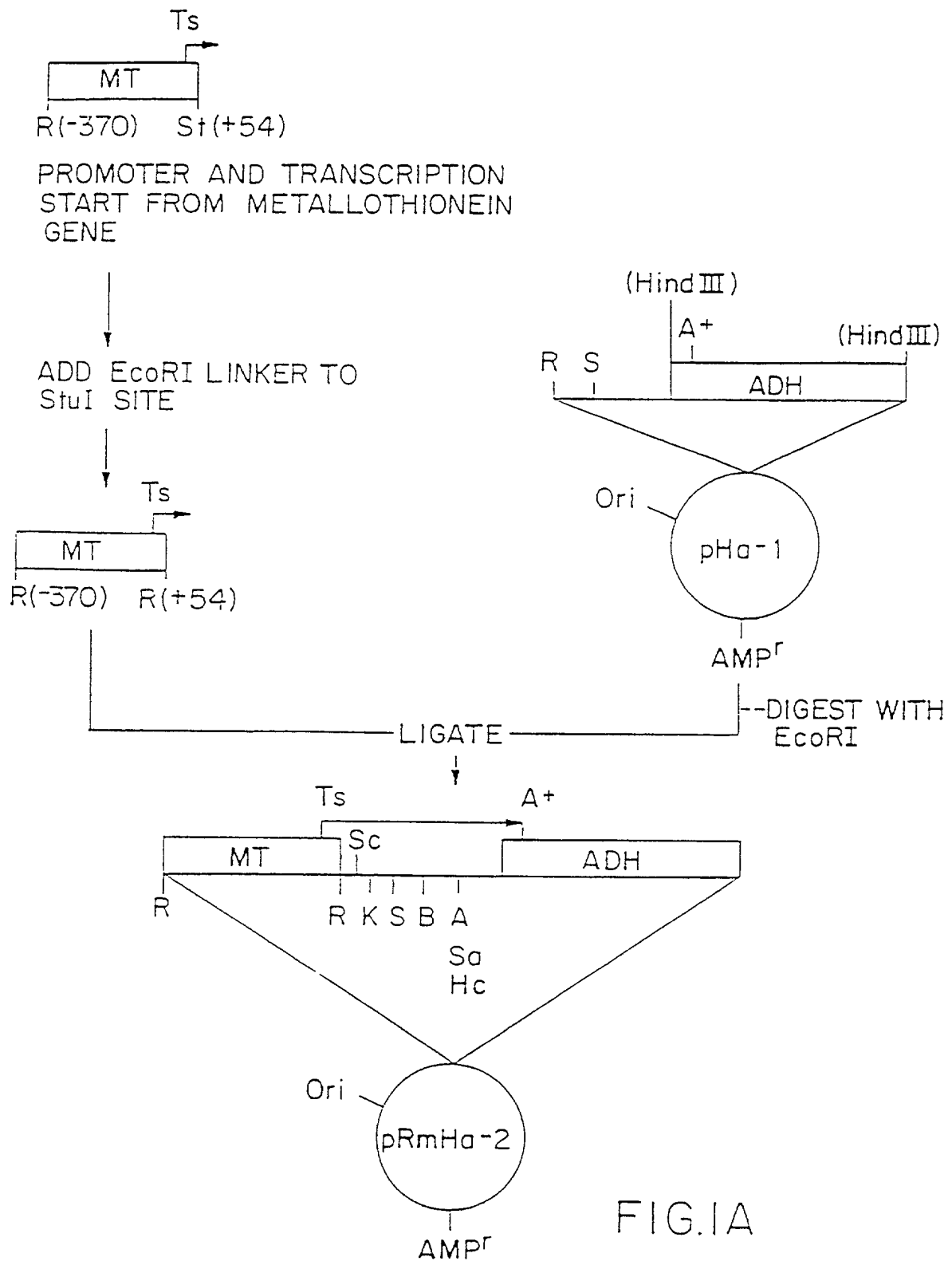
109. The method of claim 108 wherein the patient-specific cytokine profile comprises the cytokine selected from the group consisting of interleukin-2, interferon- γ and tumor necrosis factor.

110. The method of claim 105 wherein the condition is an allergy.

111. The method of claim 110 wherein the allergy is selected from the group consisting of asthma and contact sensitivity.

112. The method of claim 110 wherein the patient-specific cytokine profile is produced by a CD4⁺ Th2 type response.

113. The method of claim 112 wherein the patient-specific cytokine profile comprises the cytokine selected from the group consisting of interleukin-4 and interleukin-10.



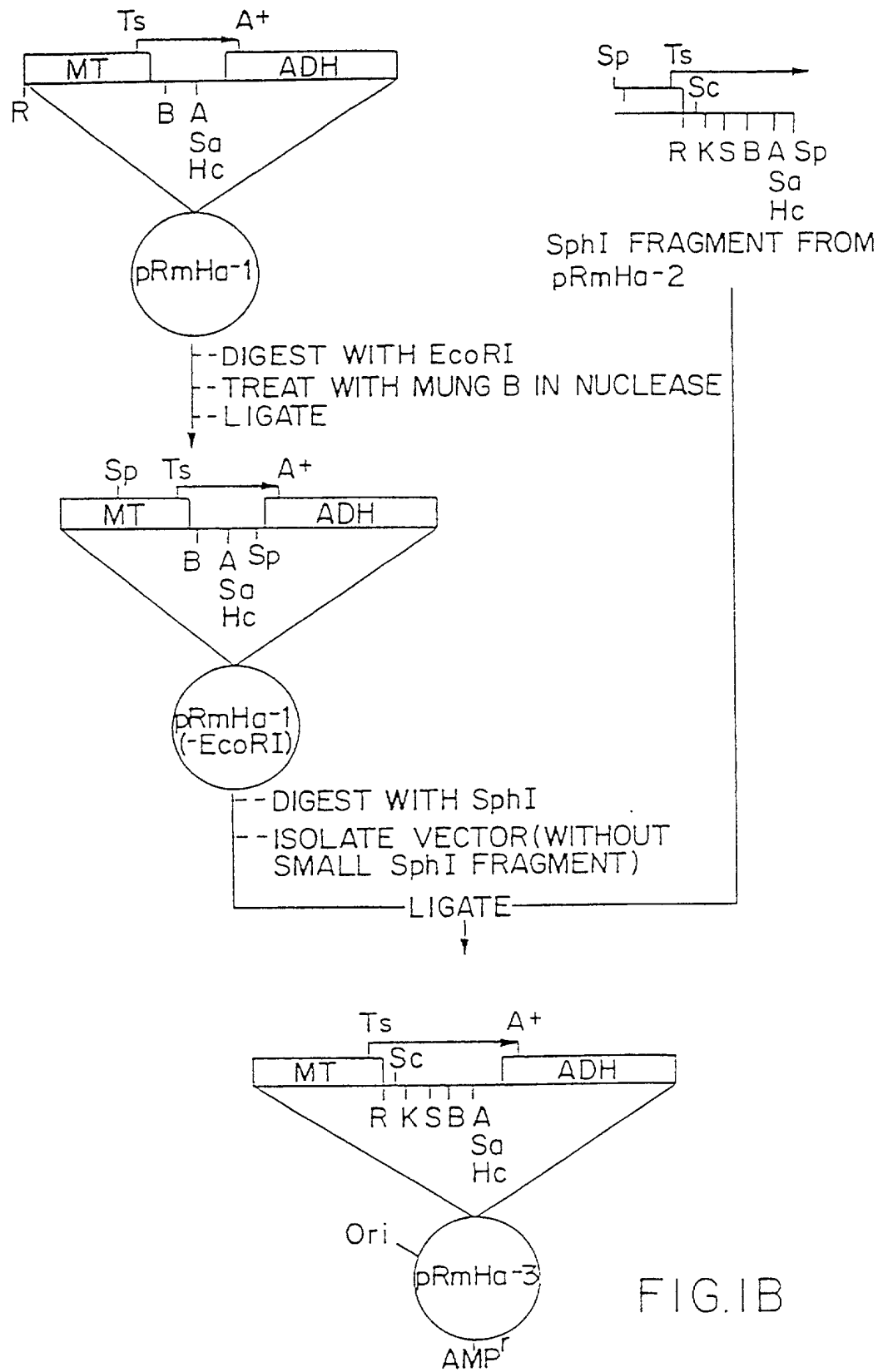


FIG.1B

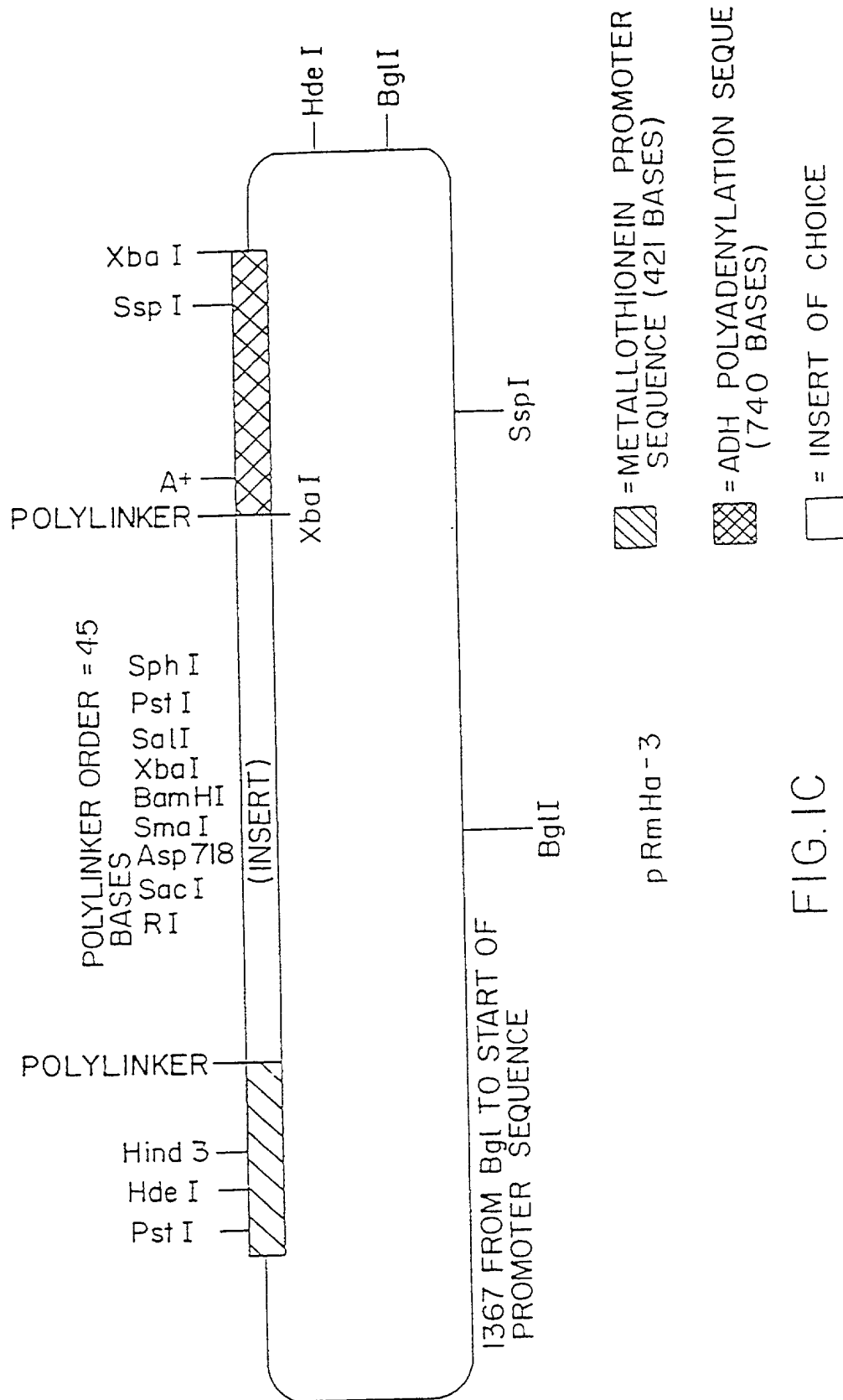


FIG. 1C

FIGURE 2

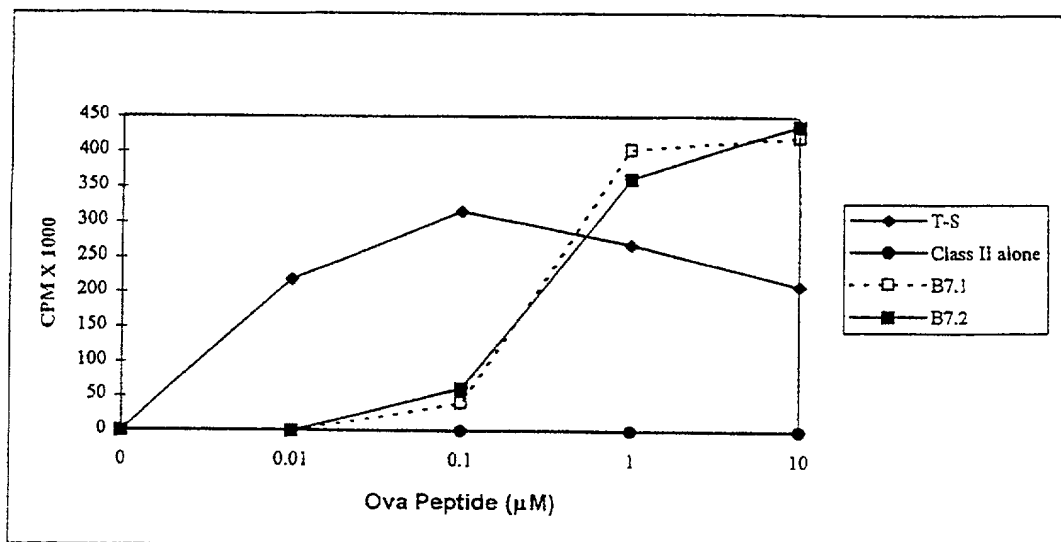
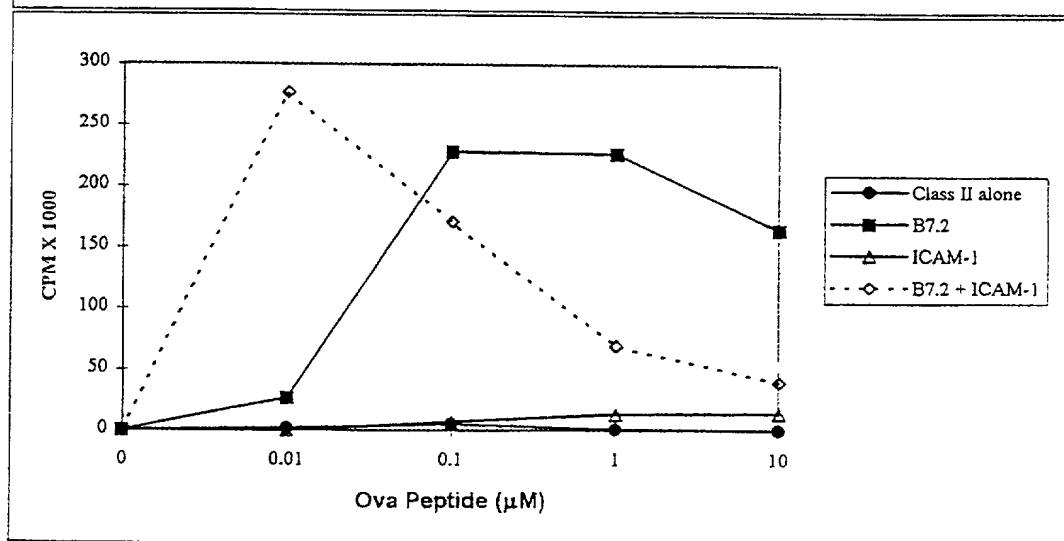


FIGURE 3



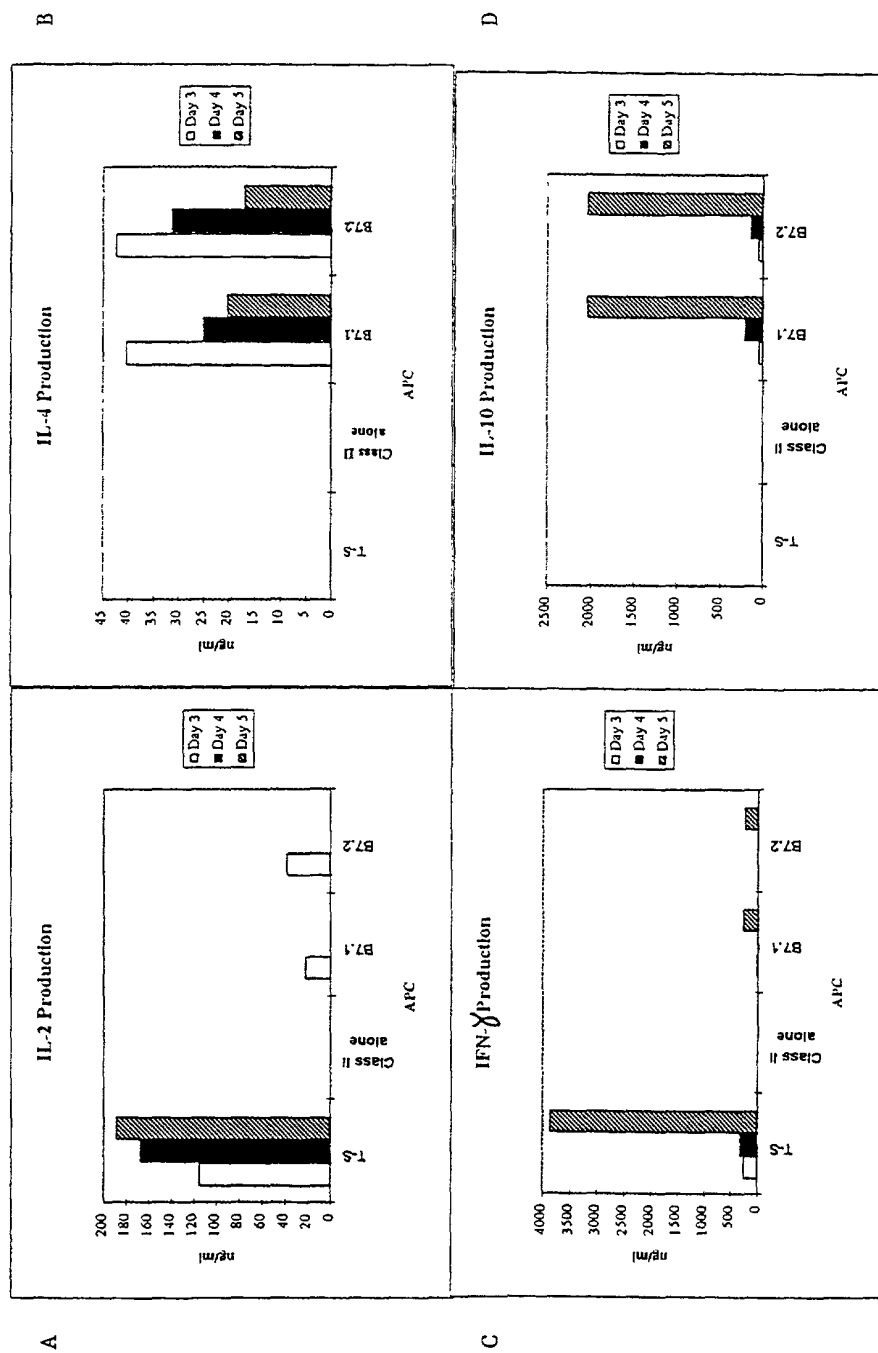


FIGURE 4

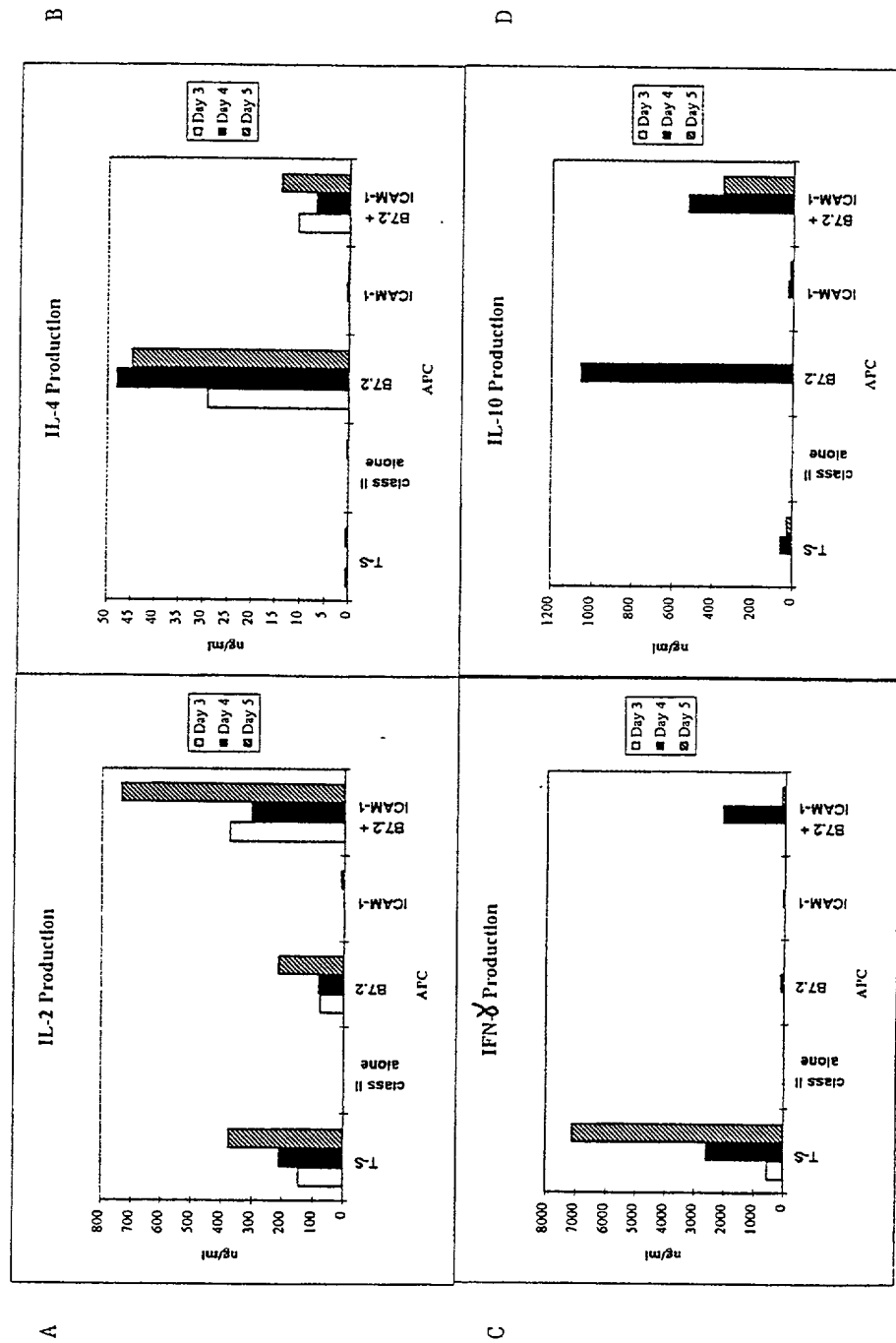


FIGURE 5

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT:

- (A) NAME: THE SCRIPPS RESEARCH INSTITUTE
- (B) STREET: 10550 North Torrey Pines Road
- (C) CITY: La Jolla
- (D) STATE: California
- (E) COUNTRY: US
- (F) ZIP: 92037
- (G) TELEPHONE: (619) 784-2937
- (H) TELEFAX: (619) 784-9399

(ii) TITLE OF INVENTION: MHC CLASS II ANTIGEN PRESENTING SYSTEMS
AND METHODS FOR ACTIVATING CD4+ T CELLS

(iii) NUMBER OF SEQUENCES: 56

(iv) COMPUTER READABLE FORM:

- (A) MEDIUM TYPE: Floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(v) CURRENT APPLICATION DATA:

- (A) APPLICATION NUMBER: PCT/US97/
- (B) FILING DATE: 22-MAY-1997
- (C) CLASSIFICATION:

(vi) PRIOR APPLICATION DATA:

- (A) APPLICATION NUMBER: US 60/018,175
- (B) FILING DATE: 23-MAY-1996

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 740 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

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TAAACAGAA GTCGTGGAAG GCGGATAGAA AGAATGTTCC ATTTGACGGG CAAGCATCTC	240
TGCTATGTGG CGGATTGCGG AGGAATTGCA CTGGAGACCA GCAAGGTTCT CATGACCAAG	300
AATATAGCGG TGTGAGTGAG CGGGAAGCTC GGTTCCTGTC CAGATCGAAC TCAAACTAG	360
TCCAGCCAGT CGCTGTCGAA ACTAATTAAG TTAATGAGTT TTTCATGTTA GTTTCGCGCT	420
GAGCAACAAT TAAGTTTATG TTTCAGTTCC GCTTAGATTT CGCTGAAGGA CTTGCCACTT	480
TCAATCAATA CTTTAGAACA AAATCAAAAC TCATTCTAAT AGCTTGGTGT TCATCTTTTT	540
TTTAAATGAT AAGCATTGTT TCGTTTATAC TTTTATATT TCGATATTAA ACCACCTATG	600
AAGTTCATTT TAATCGCCAG ATAAGCAATA TATTGTGTAA ATATTGTAT TCTTTATCAG	660
GAAATTCAGG GAGACGGGGA AGTTACTATC TACTAAAAGC GAAACAATTT CTTACAGTTT	720
TACTCTCTCT ACTCTAGAGT	740

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 427 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

AATTGCTTGC AGGACAGGAT GTGGTGCCCG ATGTGACTAG CTCTTTGCTG CAGGCCGTCC	60
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CATACATATG TGGTACGCAA GTAAGAGTGC CTGCCGATGC CCCATGTGCC CCACCAAGAG	180
TTTTGCATCC CATACAAGTC CCCAAAGTGG AGAACCGAAC CAATTCTTCG CGGGCAGAAC	240
AAAAGCTTCT GCACACGTCT CCACTCGAAT TTGGAGCCGG CCGGCGTGTG CAAAAGAGGT	300
GAATCGAACG AAAGACCCGT GTGTAAAGCC GCGTTTCCAA AATGTATAAA ACCGAGAGCA	360
TCTGGCCAAT GTGCATCAGT TGTGGTCAGC AGCAAAATCA AGTGAATCAT CTCAGTGCAA	420
CTAAAGG	427

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 35 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

CTTGAATTCC ACCATGCCGT GCAGCAGAGC TCTGA	35
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(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 29 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

TTTGGATCCT CATAAAGGCC CTGGGTGTC	29
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(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 32 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

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32

(2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

TTTGGATCCT CACTGCAGGA GCCCTGCT

28

(2) INFORMATION FOR SEQ ID NO:7:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 4713 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double

(D) TOPOLOGY: circular

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

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AGCTTCTGCA	CACGTCTCCA	CTCGAATTTG	GAGCCGGCCG	GCGTGTGCAA	AAGAGGTGAA	300
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GGCTCAGGAA	TAGCAAGTCA	GTACAGACG	GCGTTTATGA	GACCAGCTTC	CTCGTCAACC	960
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AACTGTCAGA CCAAGTTTAC TCATATATAC TTTAGATTGA TTAAAACTT CATTTTTAAT	3600
TTAAAAGGAT CTAGGTGAAG ATCCTTTTTG ATAATCTCAT GACCAAAATC CCTTAACGTG	3660
AGTTTTCGTT CCACTGAGCG TCAGACCCCG TAGAAAAGAT CAAAGGATCT TCTTGAGATC	3720
CTTTTTTTCT GCGCGTAATC TGCTGCTTGC AAACAAAAAA ACCACCGCTA CCAGCGGTGG	3780
TTTGTTTGCC GGATCAAGAG CTACCAACTC TTTTCCGAA GGTAAGTGGC TTCAGCAGAG	3840
CGCAGATACC AAATACTGTC CTTCTAGTGT AGCCGTAGTT AGGCCACCAC TTCAAGAACT	3900
CTGTAGCACC GCCTACATAC CTCGCTCTGC TAATCCTGTT ACCAGTGGCT GCTGCCAGTG	3960
GCGATAAGTC GTGTCTTACC GGGTTGGACT CAAGACGATA GTTACCGGAT AAGGCGCAGC	4020
GGTCGGGCTG AACGGGGGGT TCGTGCACAC AGCCAGCTT GGAGCGAACG ACCTACACCG	4080

AACTGAGATA CCTACAGCGT GAGCATTGAG AAAGCGCCAC GCTTCCCGAA GGGAGAAAGG 4140
CGGACAGGTA TCCGCTAAGC GGCAGGGTCG GAACAGGAGA GCGCACGAGG GAGCTTCCAG 4200
GGGGAAACGC CTGGTATCTT TATAGTCCTG TCGGGTTTCG CCACCTCTGA CTTGAGCGTC 4260
GATTTTTGTG ATGCTCGTCA GGGGGGCGGA GCCTATGGAA AAACGCCAGC AACGCGGCCT 4320
TTTTACGGTT CCTGGCCTTT TGCTGGCCTT TTGCTCACAT GTTCTTTCCT GCGTTATCCG 4380
CTGATTCTGT GGATAACCGT ATTACCGCCT TTGAGTGAGC TGATACCGCT CGCCGCAGCC 4440
GAACGACCGA GCGCAGCGAG TCAGTGAGCG AGGAAGCGGA AGAGCGCCCA ATACGCAAAC 4500
GGCTCTCCC CGCGCGTTGG CCGATTCAAT AATGCAGCTG GCACGACAGG TTTCCCGACT 4560
GGAAACGGG CAGTGAGCGC AACGCAATTA ATGTGAGTTA GCTCACTCAT TAGGCACCCC 4620
AGGCTTTACA CTTTATGCTT CCGGCTCGTA TGTGTGTGG AATTGTGAGC GGATAACAAT 4680
TTCACACAGG AAACAGCTAT GACCATGATT ACC 4713

(2) INFORMATION FOR SEQ ID NO:8:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 4724 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: circular

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

GCGTTGCAGG ACAGGATGTG GTGCCCAGTG TGA TAGCTC TTTGCTGCAG GCGGTCCTAT 60
CCTCTGGTTC CGATAAGAGA CCCAGAACTC CGGCCCCCA CGGCCACCG CCACCCCAT 120
ACATATGTGG TACGCAAGTA AGAGTGCTG CGCATGCCCC ATGTGCCCCA CCAAGAGTTT 180
TGCATCCCAT ACAAGTCCCC AAAGTGAGTA ACCGAACCAA TTCTTCGCGG GCAGAACAAA 240
AGCTTCTGCA CACGTCTCCA CTCGAATTTG GAGCCGGGCG GCGTGTGCAA AAGAGGTGAA 300
TCGAACGAAA GACCCGTGTG TAAAGCGCGG TTTCCAAAAT GTATAAAACC GAGAGCATCT 360
GGCCAATGTG CATCAGTTGT GGTGAGCAGC AAAATCAAGT GAATCATCTC AGTGCAACTA 420
AAGGGGGGAA TTCCCTGCTG TGCCCTAGAG ATGGCTCTGC AGATCCCCAG CCTCCTCCTC 480
TCAGCTGCTG TGGTGGTGCT GATGGTGCTG AGCAGCCCAG GGA CTGAGGG GGGAAACTCC 540
GAAAGGCATT TCGTGGTCCA GTTCAAGGGC GAGTGCTACT ACACCAACGG GACGCAGCGC 600
ATACGGCTCG TGACCAGATA CATCTACAAC CGGGAGGAGT ACGTGCGCTA CGACAGCGAC 660
GTGGGCGAGT ACCGCGCGGT GACCGAGCTG GGGCGGCCAG ACGCGAGTA CTGGAACAGC 720

CAGCCGGAGA TCCTGGAGCG AACGCGGGCC GAGGTGGACA CGGCGTGCAG ACACAACTAC	780
GAGGGGCGGG AGACCAGCAC CTCCCTGCCG CGGCTTGAAC AGCCCAATAT CGCCATCTCC	840
CTGTCCAGGA CAGAGGCCCT CAACCACCAC AACACTCTGG TCTGTTGGGT GACAGATTTC	900
TACCCAGCCA AGATCAAAGT GCGCTGTTT AGGAATGGCC AGGAGGAGAC AGTGGGGGTC	960
TCATCCACAC AGCTTATTAG GAATGGGGAC TGGACCTTCC AGGTCTGGT CATGCTGGAG	1020
ATGACCCCTC ATCAGGGAGA GGTCTACACC TGCCATGTGG AGCATCCCAG CCTGAAGAGC	1080
CCCATCACTG TGGAGTGGAG GGCACAGTCC GAGTCTGCCC GGAGCAAGAT GTTGAGCGGC	1140
ATCGGGGGCT GCGTGCTTGG GGTGATCTTC CTCGGGCTCG GCCTTTTCAT CCGTCACAGG	1200
ACTCAGAAAG GACCTCGAGG CCCTCCTCCA GCAGGGCTCC TGCAGTGA CTGAGTGTTC	1260
TGACTCAGTT GACTGTCTCA GACTGTAAGA CCTACATGTC TCGAATTCTGA GCTCGGTACC	1320
CGGGGATCCT CTAGAGTCGA CCTGCAGGCA TGCAATTCTGA TGCACACTCA CATTCTTCTC	1380
CTAATACGAT AATAAACTT TCCATGAAAA ATATGGAAAA ATATATGAAA ATTGAGAAAT	1440
CCAAAAAAGT GATAAACGCT CTAATTAATT AAAATAGATA AATGGGAGCG GCAGGAATCG	1500
CGGAGCATGG CCAAGTTCCT CCGCCAATCA GTCGTAAAAC AGAAGTCGTG GAAAGCGGAT	1560
AGAAAGAATG TTCGATTGA CGGGCAAGCA TGTCTGCTAT GTGGCGGATT GCGGAGGAAT	1620
TGCACTGGAG ACCAGCAAGG TTCTCATGAC CAAGAATATA GCGGTGAGTG AGCGGGAAGC	1680
TCGGTTTCTG TCCAGATCGA ACTCAAACT AGTCCAGCCA GTCGCTGTCTG AAATAATTA	1740
AGTAAATGAG TTTTTCATGT TAGTTTCGCG CTGAGCAACA ATTAAGTTTA TGTTCAGTT	1800
CGGCTTAGAT TTCGCTGAAG GACTTGCCAC TTTCAATCAA TACTTTAGAA CAAAATCAAA	1860
ACTCATTCTA ATAGCTTGGT GTTCATCTTT TTTTAAATG ATAAGCATT TGTGCTTTAT	1920
ACTTTTTATA TTTGATATT AAACCACCTA TGAAGTTCAT TTTAATCGCC AGATAAGCAA	1980
TATATTGTGT AAATATTTGT ATTCTTTATC AGGAAATTCA GGGAGACGGG GAAGTTACTA	2040
TCTACTAAAA GCCAAACAAT TTCTTACAGT TTTACTCTCT CTAATCTAGA GCTTGGCACT	2100
GGCCGTCGTT TTACAACGTC GTGACTGGGA AAACCCTGGC GTTACCCAAC TTAATCGCCT	2160
TGCAGCACAT CCCCCTTTCTG CCAGCTGGCG TAATAGCGAA GAGGCCCGCA CCGATCGCCC	2220
TTCCCAACAG TTGCGCAGCC TGAATGGCGA ATGGCGCCTG ATGCGGTATT TTCTCCTTAC	2280
GCATCTGTGC GGTATTTTAC ACCGCATATG GTGCACTCTC AGTACAATCT GCTCTGATGC	2340
CGCATAGTTA AGCCAGCCCC GACACCCGCC AACACCCGCT GACGCGCCCT GACGGGCTTG	2400
TCTGCTCCGG GCATCCGCTT ACAGACAAGC TGTGACCGTC TCCGGGAGCT GCATGTGTCA	2460
GAGGTTTTCA CCGTCATCAC CGAAACGGCG GAGACGAAAG GGCTCGTGA TACGCCTATT	2520
TTTATAGGTT AATGTCATGA TAATAATGGT TTCTTAGACG TCAGGTGGCA CTTTTCGGGG	2580
AAATGTGCGC GGAACCCCTA TTTGTTTATT TTTCTAAATA CATTCAAATA TGTATCCGCT	2640
CATGAGACAA TAACCCTGAT AAATGCTTCA ATAATATTGA AAAAGGAAGA GTATGAGTAT	2700
TCAACATTTT CGTGTCGCCC TTATTCCCTT TTTTGGGGCA TTTTGCCTTC CTGTTTTTGC	2760
TCACCCAGAA ACGCTGGTGA AAGTAAAAGA TGCTGAAGAT CAGTTGGGTG CACGAGTGGG	2820
TTACATCGAA CTGGATCTCA ACAGCGGTAA GATCCTTGAG AGTTTTCGCC CCGAAGAACG	2880
TTTTCCAATG ATGAGCACTT TTAAAGTTCT GCTATGTGGC GCGGTATTAT CCGGTATTGA	2940
CGCCGGGCAA GAGCAACTCG GTCGCGGCAT AACTATTCT CAGAATGACT TGTTGAGTA	3000
CTCACCAGTC ACAGAAAAGC ATCTTACGGA TGGCATGACA GTAAGAGAAT TATGCAGTGC	3060

TGCCATAACC	ATGAGTGATA	ACACTGCGGC	CAACTTACTT	CTGACAACGA	TCGGAGGACC	3120
GAAGGAGCTA	ACCGCTTTTT	TGCACAACAT	GGGGGATCAT	GTAACCTGCC	TTGATCGTTG	3180
GGAACCGGAG	CTGAATGAAG	CCATACCAAA	CGACGAGCGT	GACACCACGA	TGCCTGTAGC	3240
AATGGCAACA	ACGTTGCGCA	AACTATTAAC	TGGCGAACTA	CTTACTCTAG	CTTCCCGGCA	3300
ACAATTAATA	GA CTGGATGG	AGGCGGATAA	AGTTGCAGGA	CCACTTCTGC	GCTCGGGCGT	3360
TCCGGCTGGC	TGTTTATTG	CTGATAAATC	TGGAGCCGGT	GAGCGTGGGT	CTCGCGGTAT	3420
CATTGCAGCA	CTGGGGCCAG	ATGGTAAGCC	CTCCCGTATC	G TAGTTATCT	ACACGACGGG	3480
GAGTCAGGCA	ACTATGGATG	AACGAAATAG	ACAGATCGCT	GAGATAGGTG	CCTCACTGAT	3540
TAAGCATTGG	TA ACTGTCAG	ACCAAGTTTA	CTCATATATA	CTTTAGATTG	ATTTAAAACT	3600
TCATTTTTAA	TTTAAAAGGA	TCTAGGTGAA	GATCCTTTTT	GATAATCTCA	TGACCAAAAT	3660
CCCTTAACGT	GAGTTTTCGT	TCCACTGAGC	GTCAGACCCC	G TAGAAAAAG	TCAAAGGATC	3720
TTCTTGAGAT	CCTTTTTTTC	TGCGCGTAAT	CTGCTGCTTG	CAAACAAAAA	AACCACCGCT	3780
ACCAGCGGTG	GTTTGTTCG	CGGATCAAGA	GCTACCAACT	CTTTTCCGA	AGGTAAC TGG	3840
CTTCAGCAGA	GCGCAGATAC	CAAATACTGT	CCTTCTAGTG	TAGCCGTAGT	TAGGCCACCA	3900
CTTCAAGAAC	TCTGTAGCAC	CGCCTACATA	CCTCGCTCTG	CTAATCCTGT	TACCACTGGC	3960
TGCTGCCAGT	GGCGATAAGT	CGTGTCTTAC	CGGGTTGGAC	TCAAGACGAT	AGTTACCGGA	4020
TAAGGCGCAG	CGGTGGGGCT	GAACGGGGGG	TTGGTGCACA	CAGCCCAGCT	TGGAGCGAAC	4080
GACCTACACC	GA ACTGAGAT	ACCTACAGCG	TGAGCATTGA	GAAAGCGCCA	CGCTTCCCGA	4140
AGGGAGAAAG	GCGGACAGGT	ATCCGGTAAG	CGGCAGGGTC	GGAACAGGAG	AGCGCACGAG	4200
GGAGCTTCCA	GGGGGAAACG	CCTGGTATCT	TTATAGTCCT	GTGGGGTTTC	GCCACCTCTG	4260
ACTTGAGCGT	CGATTTTTGT	GATGCTCGTC	AGGGGGGCGG	AGCCTATGGA	AAAACGCCAG	4320
CAAGCGGGCC	TTTTTACGGT	TCCTGGCCTT	TTGCTGGCCT	TTTGCTCACA	TGTTCTTTCC	4380
TGCGTTATCC	CCTGATTCTG	TGGATAACCG	TATTACCGCC	TTTGAGTGAG	CTGATACCGC	4440
TCGCCGCAGC	CGAACGACCG	AGCGCAGCGA	GTCAGTGAGC	GAGGAAGCGG	AAGAGCGCCC	4500
AATACGCAAA	CCGCCTCTCC	CCGCGCGTTG	GCGGATT CAT	TAATGCAGCT	GGCAGGACAG	4560
GTTTCCCGAC	TGAAAGCGG	GCAGTGAGCG	CAACGCAATT	AATGTGAGTT	AGCTCACTCA	4620
TTAGGCACCC	CAGGCTTTAC	ACTTTATGCT	TCCGGCTCGT	ATGTTGTGTG	GAATTGTGAG	4680
CGGATAACAA	TTTACACAG	GAAACAGCTA	TGACCATGAT	TACG		4724

(2) INFORMATION FOR SEQ ID NO:9:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 23 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

CCACCATGGC CATTAGTGA GTC

23

(2) INFORMATION FOR SEQ ID NO:10:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 29 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

TTTGGATCCT TACAGAGGCC CCCTGCGTT

29

(2) INFORMATION FOR SEQ ID NO:11:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

CCACCATGGT GTGTCTGAGG CTCC

24

(2) INFORMATION FOR SEQ ID NO:12:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 29 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

TTTGATCCT CAGCTCAGGA ATCCTCTTG

29

(2) INFORMATION FOR SEQ ID NO:13:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

GCACCATGGT CCTAAACAAA GCTCTGAT

28

(2) INFORMATION FOR SEQ ID NO:14:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

TTTGGATCCT CACAAGGGCC CTTGGTGTCT

30

(2) INFORMATION FOR SEQ ID NO:15:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

CCACCATGGC TTGAAGAAG GCCTTT

26

(2) INFORMATION FOR SEQ ID NO:16:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

TTTAGATCTC AGTGCAGAAG CCCTTT

26

(2) INFORMATION FOR SEQ ID NO:17:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

CCACCATGGG CCCTGAAGAC AGAAT

25

(2) INFORMATION FOR SEQ ID NO:18:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 27 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

TTTGGATCCT CACAGGGTCC CCTGGGC

27

(2) INFORMATION FOR SEQ ID NO:19:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 26 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

CCACCATGGT TCTGCAGGT TCTGCG

26

(2) INFORMATION FOR SEQ ID NO:20:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 29 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

TTTGATCCT TATGCAGATC CTCGTTGAA

29

(2) INFORMATION FOR SEQ ID NO:21:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 26 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

AAGAATTCAC TAGAGGCTAG AGCCAT

26

(2) INFORMATION FOR SEQ ID NO:22:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

AAGGATCCTC ACAGGGTGAC TTGACC

26

(2) INFORMATION FOR SEQ ID NO:23:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 2580 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: double

(D) TOPOLOGY: circular

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

GCGTTGCAGG ACAGGATGTG GTGCCGGATG TGA CTAGCTC TTTGCTGCAG GCCGTCCTAT	60
CCTCTGGTTC CGATAAGAGA CCCAGAACTC CGGCCCCCA CCGCCCACCG CCACCCCAT	120
ACATATGTGG TACGCAAGTA AGAGTGCTG CGCATGCCCC ATGTGCCCCA CCAAGAGTTT	180
TGCATCCCAT ACAAGTCCCC AAAGTGGAGA ACCGAACCAA TTCTTCGCGG GCAGAACAAA	240
AGCTTCTGCA CAGGTCTCCA CTCGAATTTG GAGCCGGCCG GCCTGTGCAA AAGAGGTGAA	300
TCGAACGAAA GACCCGTGTG TAAAGCCGCG TTTCCAAAAT GTATAAAACC GAGAGCATCT	360
GGCCAATGTG CATCAGTTGT GGTGAGCAGC AAAATCAAGT GAATCATCTC AGTGCAACTA	420
AAGGGGGGAA TTCGATCTAG AGGCTAGAGC CATGGATGAC CAACGCGACC TCATCTCTAA	480
CCATGAGCAA TTGCCCATAC TGGGCAACCG CCCTAGAGAG CCAGAAAGGT GCAGCCGTGG	540
AGCTCTGTAC ACCGGTGTTC CTGTCCTGGT GGCTCTGCTC TTGGCTGGGC AGGCCACCAC	600
TGCTTACTTC CTGTACCAGC AACAGGGCCG CCTAGACAAG CTGACCATCA CCTCCGAGAA	660
CCTGCAACTG GAGAGCCTTC GCATGAAGCT TCGGAAATCT GCCAAACCTG TGAGCCAGAT	720
GCGGATGGCT ACTCCCTTGC TGATGCGTCC AATGTCCATG GATAACATGC TCCTTGGGCC	780
TGTGAAGAAC GTTACCAAGT ACGGCAACAT GACCCAGGAC CATGTGATGC ATCTGCTCAC	840
GAGGTCTGGA CCCCTGGAGT ACCCGCAGCT GAAGGGGACC TTCCCAGAGA ATCTGAAGCA	900
TCTTAAGAAC TCCATGGATG GCGTGAAGT GAAGATCTTC GAGAGCTGGA TGAAGCACTG	960
GCTCTTGTTC GAGATGAGCA AGAACTCCCT GGAGGAGAAG AAGCCACAG AGGCTCCACC	1020
TAAAGAGCCA CTGGACATGG AAGACCTATC TTCTGGCCTG GGAGTGACCA GGCAGGAACT	1080
GGGTCAAGTC ACCCTGTGAA GACAGAGGCC AGCATCAAGC TTATCGATAC CGTCGACCTG	1140
CAGGCATGCA ATTCGATGCA CACTCACATT CTCTCTCTAA TACGATAATA AAAGTTTCCA	1200
TGAAAAATAT GAAAAATAT ATGAAATTG AGAAATCCAA AAAAGTATA AACGCTCTAC	1260
TTAATTAAAA TAGATAAATG GGAGCGGCAG GAATGGCGGA GCATGGCCAA GTTCTCTCGC	1320
CAATCAGTCG TAAACAGAA GTCGTGAAA GCGGATAGAA AGAATGTTTC ATTTGACGGG	1380
CAAGCATGTC TGCTATGTGG CGGATTGCGG AGGAATTGCA CTGAGACCA GCAAGGTTCT	1440
CATGACCAAG AATATAGCGG TGAGTGAGCG GGAAGCTCGG TTTCTGTCCA GATCGAACTC	1500
AAAACTAGTC CAGCCAGTCG CTGTCGAAAC TAATTAAGTA AATGAGTTTT TCATGTAGT	1560
TTGCGCTGA GCAACAATTA AGTTTATGTT TCAGTTTCGGC TTAGATTTTC CTGAAGGACT	1620
TGCCACTTTC AATCAATACT TTAGAACAAA ATCAAACTC ATTCTAATAG CTGCGTGTTC	1680
ATCTTTTTTT TTAATGATAA GCATTTTGTG GTTTATACTT TTTATATTTT GATATTAAAC	1740
CACCTATGAA GTTCATTTTA ATCGCCAGAT AAGCAATATA TTGTGTAAAT ATTTGTATTC	1800
TTTATCAGGA AATTCAGGGA GACGGGGAAG TTAATATCTA CTAAGGCCA AACAATTTCT	1860
TACAGTTTTA CTCTCTCTAC TCTAGAGCTT GGCAGTGGCC GTCGTTTTAC AACGTCGTGA	1920
CTGGGAAAAC CCTGGCGTTA CCGAACTTAA TCGCCTTGCA GCACATCCCC CTTTCGCCAG	1980
CTGGCGTAAT AGCGAAGAGG CCCGCACCGA TCGCCCTTCC CAACAGTTGC GCAGCCTGAA	2040
TGGCGAATGG CGCCTGATGC GGTATTTTCT CCTTACGCAT CTGTGCGGTA TTTCACACCG	2100

CATATGGTGC	ACTCTCAGTA	CAATCTGCTC	TGATGCCGCA	TAGTTAAGCC	AGCCCCGACA	2160
CCCCCAACA	CCCGCTGACG	CGCCCTGACG	GGCTTGTCTG	CTCCGGGCAT	CCGCTTACAG	2220
ACAAGCTGTG	ACCGTCTCCG	GGAGCTGCAT	GTGTCAGAGG	TTTTCACCGT	CATCACCGAA	2280
ACGCGCGAGA	CGAAAGGGCC	TCGTGATACG	CCTATTTTTA	TAGGTTAATG	TCATGATAAT	2340
AATGGTTTCT	TAGACGTCAG	GTGGCACTTT	TCGGGGAAAT	GTGCGCGGAA	CCCCTATTTG	2400
TTTATTTTTTC	TAAATACATT	CAAATATGTA	TCCGCTCATG	AGACAATAAC	CCTGATAAAT	2460
GCTTCAATAA	TATTGAAAAA	GGAAGAGTAT	GAGTATTCAA	CATTTCCGTG	TCGCCCTTAT	2520
TCCCTTTTTTT	GCGGCATTTT	GCCTTCCTGT	TTTTGCTCAC	CCAGAAACGC	TGGTGAAAGT	2580

(2) INFORMATION FOR SEQ ID NO:24:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 32 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

AAGAATTCAC CATGGATGAT CAGCGCGACC TT

32

(2) INFORMATION FOR SEQ ID NO:25:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 31 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

AAAGGATCCT CACATGGGGA CTGGGCCAG A

31

(2) INFORMATION FOR SEQ ID NO:26:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 25 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

AAACCATGGG TCATGAACAG AACCA

25

(2) INFORMATION FOR SEQ ID NO:27:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 27 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

TTTGTGCACT CAGTCACCTG AGCAAGG

27

(2) INFORMATION FOR SEQ ID NO:28:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 22 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

AAACCATGGT CTCATTCCTG CC

22

(2) INFORMATION FOR SEQ ID NO:29:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:

TTTGTGGACC TAGGAAATGT GCCATCC

27

(2) INFORMATION FOR SEQ ID NO:30:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 34 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

TTTAGAATTC ACCATGGCTT CAACCCCTGC CAAG

34

(2) INFORMATION FOR SEQ ID NO:31:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:

TTTAGTCGAC TCAGGGAGGT GGGGCTTGTC C

31

(2) INFORMATION FOR SEQ ID NO:32:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 36 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:

ACCCTTGAAT TCATGGCTCC CAGCAGCCCC CGGCCC

36

(2) INFORMATION FOR SEQ ID NO:33:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 39 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:

ATTACCGGAT CCTCAGGGAG GCGTGGCTTG TGTGTTCCG

39

(2) INFORMATION FOR SEQ ID NO:34:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 27 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:

AAGGTACCCG TGGAGACTGC CAGAGAT

27

(2) INFORMATION FOR SEQ ID NO:35:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:

TTTGGATCCC TATGGCCGGA AGGCCTG

27

(2) INFORMATION FOR SEQ ID NO:36:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:

AAGAATTCCT GTCAGAATGG CCACCAT

27

(2) INFORMATION FOR SEQ ID NO:37:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:

TTTAGATCTT CACTCAGCTC TGGACGGT

28

(2) INFORMATION FOR SEQ ID NO:38:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 36 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

ACCCTTGAGC TCATGGTTGC TGGGAGCGAC GCGGGG

36

(2) INFORMATION FOR SEQ ID NO:39:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 42 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:

ATTACCGGAT CCTTAAAGAA CATTATATA CAGCACAATA CA

42

(2) INFORMATION FOR SEQ ID NO:40:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 34 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:

TTTAGAATTC ACCATGGCTT GCAATTGTCA GTTG

34

(2) INFORMATION FOR SEQ ID NO:41:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:

TTTAGTCGAC CTAAAGGAAG ACGGTCTGTT C

31

(2) INFORMATION FOR SEQ ID NO:42:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:

ACCCTTGAAT CCATGGGCCA CACACGGAGG CAG

33

(2) INFORMATION FOR SEQ ID NO:43:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 39 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:

ATTACCGGAT CCTTATACAG GCGGTACACT TTCCCTTCT

39

(2) INFORMATION FOR SEQ ID NO:44:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 36 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:

TTTAGAATTC ACCATGGACC CCAGATGCAC GATGGG

36

(2) INFORMATION FOR SEQ ID NO:45:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 34 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:

TTTAGTCGAC TCACTCTGCA TTTGGTTTTG CTGA

34

(2) INFORMATION FOR SEQ ID NO:46:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 33 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:

ACCCTTGAGC TCATGGATCC CCAGTGCACT ATG

33

(2) INFORMATION FOR SEQ ID NO:47:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 42 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:

ATTACCCCCG GGTAAAAAAC ATGTATCACT TTTGTCGCAT GA

42

(2) INFORMATION FOR SEQ ID NO:48:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 31 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:48:

AAAGGATCCA CCATGCAGCA GCCCTTCAAT T

31

(2) INFORMATION FOR SEQ ID NO:49:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 29 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:

TTTGGATCCT TAGAGCTTAT ATAAGCCGA

29

(2) INFORMATION FOR SEQ ID NO:50:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 34 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:

AAAGAATTGG GTACCATGCC GGAGGAGGCT TCGG

34

(2) INFORMATION FOR SEQ ID NO:51:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 29 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:51:

TTTGGATCCT CAGGGGGCGCA CCCACTGCA

29

(2) INFORMATION FOR SEQ ID NO:52:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 17 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:52:

Ile Ser Gln Ala Val His Ala Ala His Ala Glu Ile Asn Glu Ala Gly

1

5

10

15

Arg

(2) INFORMATION FOR SEQ ID NO:53:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 13 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:53:

Pro Lys Tyr Val Lys Gln Asn Thr Leu Lys Leu Ala Thr
1 5 10

(2) INFORMATION FOR SEQ ID NO:54:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 11 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:54:

Lys Thr Ile Ala Thr Asp Glu Glu Ala Arg Arg
1 5 10

(2) INFORMATION FOR SEQ ID NO:55:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:

Gln Ala Ser Leu Ala Leu Ser Tyr Arg Leu Asn Met Phe Thr Pro
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:56:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 13 amino acids
- (B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:

Phe	Val	Arg	Phe	Asp	Ser	Asp	Ala	Ala	Ser	Gln	Arg	Met
1				5					10			

PATENT APPLICATION DECLARATION AND POWER OF ATTORNEY

I HEREBY DECLARE THAT:

My residence, post office address, and citizenship are as stated next to my name in PART A of page 2 hereof.

I believe I am the original, first, and sole inventor (if only one name is listed) or an original, first, and joint inventor (if plural names are listed) of the subject matter which is claimed and for which a patent is sought on the invention entitled MHC CLASS II ANTIGEN PRESENTING SYSTEMS AND METHODS FOR ACTIVATING CD4+ T CELLS the specification of which:

_____ is attached hereto
X was filed on May 22, 1997 as Application Serial No. PCT/US97/08697
and was amended on _____ (if applicable).

I hereby state that I have reviewed and understand the contents of the above-identified specification, including the claims, as amended by any amendment referred to above.

I acknowledge the duty to disclose information which is material to the examination of the application in accordance with Title 37, Code of Federal Regulations, Sec. 1.56(a).

I hereby claim foreign priority benefits under Title 35, United States Code, Sec. 119 of any foreign application(s) for patent or inventor's certificate listed in PART B on page 2 hereof and have also identified in PART B on page 2 hereof any foreign application for patent or inventor's certificate having a filing date before that of the application on which priority is claimed.

I hereby claim the benefit under Title 35, United States Code, Sec. 120 of any United States application(s) listed in PART C on page 2 hereof and, insofar as the subject matter of each of the claims of this application is not disclosed in the prior United States application in the manner provided by the first paragraph of Title 35, United States Code, Sec. 112, I acknowledge the duty to disclose material information as defined in Title 37, Code of Federal Regulations, Sec. 1.56(a) which occurred between the filing date of the prior application and the national or PCT international filing date of this application.

I hereby declare that all statements made herein of my knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

I hereby appoint the following as my attorneys or agents with full power of substitution to prosecute this application and transact all business in the United States Patent and Trademark Office connected therewith:

Douglas A. Bingham	Reg. No. 32,457	Emily Holmes	Reg. No. 40,652
Thomas Fitting	Reg. No. 34,163	Thomas E. Northrup	Reg. No. 33,268
Donald G. Lewis	Reg. No. 28,636		

whose mailing address for this application is:
THE SCRIPPS RESEARCH INSTITUTE
10550 North Torrey Pines Road, Mail Drop TPC-8
La Jolla, California 92037

See Page 2 attached, signed, and made a part hereof.

PATENT APPLICATION DECLARATION AND POWER OF ATTORNEY

PART A: Inventor Information And Signature

Full name of SOLE or FIRST inventor Susan R. Webb
Citizenship USA Post Office Address 10550 N. Torrey Pines Road,
Department of Immunology, IMM4, La Jolla, CA 92037
Residence (if different) 1076 Hymettus Ave., Leucadia, CA 92024

Inventor's Signature: *Susan R. Webb* Date: 4/9/99

Full name of SECOND joint inventor, if any Ola Wingvist
Citizenship Sweden Post Office Address _____
Residence (if different) _____

Second Inventor's Signature: _____ Date: _____

Full name of THIRD joint inventor, if any Lars Karlsson
Citizenship Sweden Post Office Address _____
Residence (if different) _____

Third Inventor's Signature: _____ Date: _____

Full name of FOURTH joint inventor, if any Michael R. Jackson
Citizenship UK Post Office Address _____
Residence (if different) _____

Fourth Inventor's Signature: _____ Date: _____

Full name of FIFTH joint inventor, if any Per A. Peterson
Citizenship Sweden Post Office Address _____
Residence (if different) _____

Fifth Inventor's Signature: _____ Date: _____

Full name of SIXTH joint inventor, if any _____
Citizenship _____ Post Office Address _____
Residence (if different) _____

Sixth Inventor's Signature: _____ Date: _____

PART B: Prior Foreign Application(s)
Serial No. Country Day/Month/Year Filed Priority Claimed
____ Yes ____ No

PART C: Claim For Benefit of Filing Date of Earlier U.S. Application(s)

Serial No.	Filing Date	Status:
60/018,175	5/23/96	<input type="checkbox"/> Patented <input type="checkbox"/> Pending <input checked="" type="checkbox"/> Abandoned
		<input type="checkbox"/> Patented <input type="checkbox"/> Pending <input type="checkbox"/> Abandoned

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Thomas Fitting	Reg. No. 34,163	Thomas E. Northrup	Reg. No. 33,268
Donald G. Lewis	Reg. No. 28,636		

whose mailing address for this application is:

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La Jolla, California 92037

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Full name of SOLE or FIRST inventor Susan R. Webb
Citizenship USA Post Office Address _____

Residence (if different) _____

Inventor's Signature: _____ Date: _____

Full name of SECOND joint inventor, if any Ola Wingvist
Citizenship Sweden Post Office Address Luthagsespl. 9B
S-751 85 Uppsala, SWEDEN

Residence (if different) _____

Second Inventor's Signature: [Signature] Date: 990320

Full name of THIRD joint inventor, if any Lars Karlsson
Citizenship Sweden Post Office Address _____

Residence (if different) _____

Third Inventor's Signature: _____ Date: _____

Full name of FOURTH joint inventor, if any Michael R. Jackson
Citizenship UK Post Office Address _____

Residence (if different) _____

Fourth Inventor's Signature: _____ Date: _____

Full name of FIFTH joint inventor, if any Per A. Peterson
Citizenship Sweden Post Office Address _____

Residence (if different) _____

Fifth Inventor's Signature: _____ Date: _____

Full name of SIXTH joint inventor, if any _____
Citizenship _____ Post Office Address _____

Residence (if different) _____

Sixth Inventor's Signature: _____ Date: _____

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Serial No.	Country	Day/Month/Year Filed	Priority Claimed
			<input type="checkbox"/> Yes <input type="checkbox"/> No

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Full name of SOLE or FIRST inventor Susan R. Webb
Citizenship USA Post Office Address Work: The Scripps Research Institute, IMM-4,
10555 North Torrey Pines Road, La Jolla, CA 92037

Residence (if different) _____

Inventor's Signature: _____ Date: _____

Full name of SECOND joint inventor, if any Ola Wingqvist
Citizenship Sweden Post Office Address Work: Department of Internal Medicine,
Uppsala University Hospital, S-751 85, Uppsala, Sweden

Residence (if different) _____

Second Inventor's Signature: _____ Date: _____

Full name of THIRD joint inventor, if any Lars Karlsson
Citizenship Sweden Post Office Address Work: RWJPRI, 3535 General Atomics Court,
Suite 100, San Diego, CA 92121

Residence (if different) 554 Rosemont Street, La Jolla, CA 92037, U.S.

Third Inventor's Signature: [Signature] Date: 3/23/99

Full name of FOURTH joint inventor, if any Michael R. Jackson
Citizenship UK Post Office Address Work: RWJPRI, 3535 General Atomics Court,
Suite 100, San Diego, CA 92121

Residence (if different) 12728 Via Felino, Del Mar, CA 92014

Fourth Inventor's Signature: [Signature] Date: 3/22/99

Full name of FIFTH joint inventor, if any Per A. Peterson
Citizenship Sweden Post Office Address Work: RWJPRI, 3535 General Atomics Court,
Suite 100, San Diego, CA 92121

Residence (if different) P.O. Box 2323 Rancho Santa Fe, CA 92067

Fifth Inventor's Signature: [Signature] Date: 3/23/99

Full name of SIXTH joint inventor, if any _____
Citizenship _____ Post Office Address _____

Residence (if different) _____

Sixth Inventor's Signature: _____ Date: _____

PART B: Prior Foreign Application(s)

Serial No.	Country	Day/Month/Year Filed	Priority Claimed
			<input type="checkbox"/> Yes <input type="checkbox"/> No

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Serial No.	Filing Date	Status:
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		<input type="checkbox"/> Patented <input type="checkbox"/> Pending <input type="checkbox"/> Abandoned

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